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| <b>(54) Title:</b> USE OF 7 ALPHA-SUBSTITUTED STEROIDS TO TREAT NEUROPSYCHIATRIC, IMMUNE OR ENDOCRINE DISORDERS  |           |  |
| <b>(57) Abstract</b><br><p>Use is provided for a 7<math>\alpha</math>-hydroxy or 7-oxo substituted 3<math>\beta</math>-hydroxy-steroid possessing the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol, or an analogue thereof substituted independently at one or both of the 7- and 3-positions with an ester or ether group, in the manufacture of a pharmaceutical composition for the therapy of neuropsychiatric, immune and/or endocrine disorders or for inducing cognitive enhancement. Uses for Cyp7b enzymes in producing such steroids is also provided together with various novel steroids and test kits and methods for diagnosing the disorders.</p>  |           |  |

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## USE OF 7 ALPHA-SUBSTITUTED STEROIDS TO TREAT NEUROPSYCHIATRIC, IMMUNE OR ENDOCRINE DISORDERS

The present invention relates to novel uses for 7 $\alpha$ -hydroxy-substituted steroids, to a process for preparing such steroids and to novel steroids so produced.

In particular the invention relates to the use of cytochromes of the cytochrome  
5 P450 family designated Cyp7b to effect 7 $\alpha$ -hydroxylation of certain 3 $\beta$ -OH steroids so as to produce a 7 $\alpha$ -hydroxy-substituted steroids. Certain of the 7 $\alpha$ -hydroxy-substituted steroids so produced, as well the corresponding 7-oxo derivatives, are novel and form further aspects of the invention. The invention also relates to uses of these steroids, to uses of Cyp7b enzymes and to uses of novel macromolecular species, eg. antibodies and DNAs,  
10 which are biologically related to the Cyp7b enzymes.

Cytochromes P450 are a diverse group of heme-containing mono-oxygenases (termed CYP's; see Nelson *et al.*, DNA Cell Biol. (1993) 12, 1-51) that catalyse a variety of oxidative conversions, notably of steroids but also of fatty acids and xenobiotics. While CYP's are most abundantly expressed in the testis, ovary, placenta, adrenal and liver, it is  
15 becoming clear that the brain is a further site of CYP expression. Several CYP activities or mRNA's have been reported in the nervous system but these are predominantly of types metabolizing fatty acids and xenobiotics (subclasses CYP2C, 2D, 2E and 4). However, primary rat brain-derived glial cells have the capacity to synthesize pregnenolone and progesterone *in vitro*. Mellon and Deschepper, Brain Res. (1993), 629, 283-292(9)  
20 provided molecular evidence for the presence, in brain, of key steroidogenic enzymes CYP11A1 (sc) and CYP11B1 (11 $\beta$ ) but failed to detect CYP17 (c17) or CYP11B2 (AS). Although CYP21A1 (c21) activity is reported to be present in brain, authentic CYP21A1 transcripts were not detected in this tissue.

Interest in steroid metabolism in brain has been fuelled by the finding that adrenal-  
25 and brain-derived steroids (neurosteroids) can modulate cognitive function and synaptic plasticity. For instance, pregnenolone and steroids derived from it are reported to have memory enhancing effects in mice. However, the full spectrum of steroid metabolizing CYP's in brain and the biological roles of their metabolites *in vivo* has not been established.

Many aspects of brain function are modulated by steroids. Intracellular receptors  
30 for glucocorticoids (cortisol, corticosterone) are particularly abundantly expressed in the

hippocampus (1), a brain region that plays a key role in specific aspects of memory formation, and which is an early and prominent target for dysfunction and damage in Alzheimer's disease (AD). While glucocorticoids regulate learning and memory, mood and neuroendocrine control, chronic glucocorticoid excess compromises neuronal activity, synaptic plasticity and eventually survival, particularly in the hippocampus. These findings prompted the suggestion that glucocorticoid-mediated neurotoxicity might underpin some age-related brain disorders, including AD, in which plasma cortisol levels are markedly elevated (2).

Conversely, dehydroepiandrosterone (DHEA), the most abundant steroid product of the human adrenal cortex, has been proposed to protect against disorders of the aging brain (3). Plasma levels of DHEA often show a striking age-associated decline which correlates with loss of cognitive function (4). In rodents, injection of DHEA or its sulfate into limbic structures improves post-training memory and enhances synaptic plasticity (5). DHEA and glucocorticoids thereby appear to exert inverse effects upon memory function and synaptic plasticity, and DHEA has been advocated as an endogenous 'anti-glucocorticoid'. However, despite considerable circumstantial evidence to support this contention, there is no evidence for a direct interaction between DHEA and glucocorticoid signalling pathways in neurons.

Neurosteroidogenesis has been reported in isolated rat retina (8) and brain (9). In addition to the production of pregnenolone and DHEA from cholesterol, a variety of novel steroids are made in brain extracts or cultured brain cells, including  $20\alpha$ -dehydropregnenolone,  $7\alpha$ -hydroxy derivatives of pregnenolone and DHEA, progesterone, and both  $3\alpha$ - and  $3\beta$ -hydroxy- $5\alpha$ -pregnan-20-one (reviewed in Ref. 7). Androgens are also modified, particularly through the action of aromatase and a  $5\alpha$ -reductase (reviewed in Ref. 10). However, the specific enzymes responsible for these and other transformations in the central nervous system have not been well characterized.

As referred to above, several Cyps are present in the central nervous system (11-22). Activities or mRNAs corresponding to key steroidogenic enzymes (23-25), in addition to Cyp19 (aromatase) have been detected. Furthermore, mRNAs encoding the non-Cyp hydroxysteroid dehydrogenases (HSD)  $3\alpha$ -HSD,  $3\beta$ -HSD and  $11\beta$ -HSD have been reported in the central nervous system (25, 27-29).

To investigate regulation of brain function, studies reported in copending International Patent Application No PCT/GB95/02465, published as WO 96/12810, and in Stapleton *et al* (J. Biol. Chem. 270, 29739 - 1995, December, 15 1995), focused on the hippocampus, a brain region important in learning and memory. A copy of the specification  
5 of International Patent Application No PCT/GB95/02465 has been filed with the priority documents filed in respect of this specification.

That copending application, PCT/GB95/02465, describes and claims novel cytochrome P450 proteins designated Hct-1. These Hct-1 proteins have now been named as Cyp7b by the Committee on Standardized Cytochrome P450 Nomenclature and the  
10 name Cyp7b will be used in this application.

The Cyp7b enzyme shares 39% sequence identity to hepatic cholesterol 7 $\alpha$ -hydroxylase (Cyp7a) and lesser but significant homology with other steroidogenic Cyps. The postulated steroidogenic domain (30,31), found in many of these enzymes, is present in both Cyp7a and Cyp7b. Cyp7b mRNA is predominantly expressed in rodent brain,  
15 particularly in the hippocampus, unlike Cyp7a, which is liver-specific (31-33 and EP0648840 A2).

The present inventors have now investigated the substrate specificity of Cyp7b and found that Cyp7b catalyses the introduction of a hydroxyl group at the 7 $\alpha$  position in steroid substrates, particularly 3 $\beta$ -hydroxy steroids. Cytochromes Cyp7b are thus steroid  
20 hydroxylase enzymes having 7 $\alpha$ -specificity. The ability to produce 7 $\alpha$ -hydroxylated steroids is of major commercial importance, because such steroids are of particular use in the manufacture of pharmaceuticals (either as drugs *per se* or as intermediates), and in the manufacture of test kits and assays for pathological conditions associated with the presence of abnormal levels of endogenous enzyme, substrate or product.

25 The abbreviation "DHEA" will be used herein to designate dehydroepiandrosterone, thus 7 $\alpha$ -hydroxy-DHEA designates 7 $\alpha$ -hydroxydehydroepi-androsterone

The present inventors have identified substrate/product pairs associated with Cyp7b, particularly DHEA/7 $\alpha$ -hydroxy-DHEA (7-HD), pregnenolone/7 $\alpha$ -hydroxy-pregnenolone (7-HP) and  $\beta$ -estradiol/7 $\alpha$ -hydroxy- $\beta$ -estradiol (7-HE). They have also determined that  
30 DHEA concentration in brain tissue declines with age, whereas the concentrations of other brain steroids do not, and determined that the ageing process may be associated with

deficits in certain steroids and also with deficits in the concentration of Cyp7b itself. It is also believed that one of the products produced by Cyp7b mediated reactions, namely 7 $\alpha$ -hydroxy dehydroepiandrosterone, plays an important role in the operation of the immune system. Because 7 $\alpha$ -hydroxy-DHEA is believed to be made substantially only in the brain, the inventors hypothesize that senescence may be due to a deficit in brain-produced 7 $\alpha$ -hydroxy-DHEA as well as in other steroids found in the brain such as DHEA, pregnenolone and 7 $\alpha$ -hydroxy-pregnenolone.

The present inventors have now further determined that one of the specific properties of the 7 $\alpha$ -hydroxy-substituted steroids, and potentially their 7-oxo substituted steroid derivatives, provided by the present invention is that of glucocorticoid and/or mineralocorticoid antagonism, whether at receptor level or otherwise. This is particularly demonstrated by the Example 5 below with respect to 7 $\alpha$ -hydroxy-DHEA but is more generally applicable. Thus this activity not only gives further uses for the novel steroids of the invention but provides first and second medical uses for known 7 $\alpha$ -hydroxy or 7-oxo steroids made available by the present process as glucocorticoid and/or mineralocorticoid antagonists and preferably in antagonism specific to neuronal tissue such as in the CNS.

Thus, having regard to this activity and their involvement in endogenous metabolic pathways, particularly in the brain, the 7 $\alpha$ -hydroxy substituted 3 $\beta$ -hydroxy-steroids provided by use of the Cyp7b enzyme activity, including novel compounds provided by the invention, and their 7-oxo derivatives, have utility in the therapy of neuropsychiatric, immune and endocrine disorders, particularly but not exclusively steroid associated disorders.

Use of these 7 $\alpha$ -hydroxy or 7-oxo substituted 3 $\beta$ -hydroxy-steroids, preferably possessing the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol, or derivatives thereof substituted independently at one or both of the 7- and 3- positions with an ester or ether group, in treating these disorders and for manufacturing medicaments for such treatment is provided in a first aspect of the present invention. Particularly preferred derivatives are those wherein one or both of the ester and or ether group is metabolisable *in vivo* to produce the corresponding hydroxy compound.

Preferred derivatives include those wherein the steroid has a 3 $\beta$ -substituent-OR<sub>1</sub> and/or a 7 $\alpha$ -substituent -OR<sub>2</sub> where -OR<sub>1</sub> and -OR<sub>2</sub> each independently represents a free

hydroxy, ester or ether group,

wherein each of  $R_1$  and  $R_2$  are independently selected from the group consisting of hydrogen, substituted or unsubstituted  $C_{1-6}$  alkyl groups, groups  $R_5CO-$ , wherein  $R_5$  may be selected from substituted or unsubstituted  $C_{1-6}$  alkyl groups, and groups of the formula  $-OP(OH)_3$ , wherein any substituents are selected from OH, halogen (F, Cl, Br, I) amino,  $C_{1-6}$  alkylamino,  $C_{1-6}$  dialkylamino,  $COOH$  or  $COOR_4$  wherein  $R_4$  represents a  $C_{1-6}$  alkyl group; and wherein the compounds may be in free form or in the form of acid addition salts with pharmacologically acceptable anions.

The particular disorders for which this utility is provided include

- 10 (a) deficits of cognition in aging
- (b) Alzheimer's disease
- (c) deficits of immune system in aging
- (d) deficits of immune function in HIV infection
- (e) glucocorticoid or mineralocorticoid excess
- 15 (f) diabetes
- (g) depression
- (h) osteoporosis and hypercalcemia
- (i) hyperglycemia and hyperlipodemia
- (j) muscle atrophy
- 20 (k) arterosclerosis
- (l) steroid diabetes

Further, these  $7\alpha$ -hydroxy steroids, their esters, ethers and 7-oxo derivatives may be used to induce cognitive enhancement in a normal individual.

Preferred steroids for such use have the carbon skeleton of androsterone, pregnenolone or estradiol and particularly preferred examples are  $7\alpha$ -hydroxy-DHEA and  $7\alpha$ -hydroxypregnenolone. Accordingly the present invention further provides the use of novel compounds of Formula Ia and Ib shown below in the applications indicated above.

Particularly preferred uses for the antagonistic properties of these 7-substituted steroids include treatment of disorders falling within category (e) above or where reversal of the effects of such corticoids, regardless of excess, is required.

A second aspect of the present invention provides pharmaceutical compositions implementing such use. The compositions in which the novel steroids and known steroids of the invention will be used will readily occur to those skilled in the art, generally comprising the steroid active in association with a pharmaceutically acceptable carrier or diluent, with formulations for example being suitable for inhalation or for gastrointestinal (eg. oral), parenteral, topical, transdermal or transmucosal administration.

As an alternative to administering the compounds of the invention *per se*, a third aspect of the invention provides the possibility of using the gene sequences of the Cyp7b genes in gene therapy in order to compensate for a deficiency in Cyp7b enzyme. In such therapies, constructs comprising Cyp7b coding sequences can be packaged in conventional delivery systems, such as adenoviruses, vaccinia viruses, herpes viruses and liposomes and administered via a route which results in preferential targeting of a selected tissue, especially the brain. The invention further provides the possibility of using the gene sequences of the Cyp7b genes in gene therapy in order to achieve the endogenous expression of Cyp7b sequences for other purposes, e.g. in order to promote immunogenic processes. Thus for example, a vector such as a suitably modified vaccinia virus (or variant thereof) may be co-administered with a vaccine formulation so that the expressed Cyp7b sequences augment the immunogenic properties of the vaccine.

It will be realised that in the event of Cyp7b related disorders other than those involving its depletion it may be desirable to use vectors containing antisense sequences to Cyp7b effective such as to inhibit Cyp7b expression.

Macromolecules related immunologically to Cyp7b enzymes form fourth and fifth aspects of the invention and in this regard antibodies, particularly monoclonal antibodies which are capable of selectively binding Cyp7b, have utility in the diagnosis of disorders (a) to (l) referred to above. Anti-Cyp7b antibodies (including monoclonal antibodies) as well as binding molecules comprising antibody fragments may be produced by known methods and used in test kits for assays for Cyp7b enzymes.

According to a sixth aspect of the invention, there is provided a process of producing a  $7\alpha$ -hydroxy-substituted steroid which comprises subjecting a corresponding steroid substrate having no hydroxyl substituent in the 7-position to hydroxylation in the presence of a Cyp7b steroid hydroxylase enzyme.



The Cyp7b steroid hydroxylase enzyme used in the process of the invention is preferably a Cyp7b enzyme described and claimed in the above-mentioned International Patent Application No PCT/GB95/02465 (and referred to therein as Hct-1). Such enzymes include (a) ones having the precise amino acid sequences described for mouse, rat and human Cyp7b, (b) homologous enzymes from other species and (c) enzymes having amino acid sequences which differ from the sequences of enzymes included in definitions (a) and (b), but in which the capacity to catalyse the introduction of a 7 $\alpha$ -hydroxyl group is not eliminated.

The amino acid sequence of suitable Cyp7b steroid hydroxylase enzymes may be defined in terms of the DNA coding sequences disclosed in International Patent Application No PCT/GB95/02465. Thus the Cyp7b steroid hydroxylase enzyme may have a sequence encoded by DNA coding sequences of Cyp7b enzymes selected from

- (a) Coding sequences of DNA molecules comprising the coding sequence for rat Cyp7b set forth in SEQ Id No: 1,
- 15 (b) Coding sequences of DNA molecules comprising the coding sequence for mouse Cyp7b set forth in SEQ Id No: 2,
- (c) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecule defined in (a) or (b) under standard hybridization conditions defined as 2 x SSC at 65°C.
- 20 (d) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecule defined in (a), (b) or (c) under reduced stringency hybridization conditions defined as 6 x SSC at 55°C.

The sequences (a) and (b) above represent rat and mouse Hct-1 gene sequence. Homologous sequences from other vertebrate species, especially mammalian species (including man) fall within the class of DNA molecules represented by (c) or (d).

Thus for human Cyp7b, the steroid hydroxylase enzyme may comprise a sequence encoded by

- (e) DNA coding sequences selected from the following:
- (i) the sequence designated "exon 3" in SEQ Id No 3,
  - (ii) the sequence designated "exon 4" in SEQ Id No 3, and
- (f) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecules defined in (e) under standard hybridization conditions defined as 2 x SSC at 65°C.
- 
- (g) Cyp7b steroid hydroxylase encoding DNA molecules capable of hybridizing with the DNA molecule defined in (e) or (f) under reduced stringency hybridization conditions defined as 6 x SSC at 55°C.
- (h) Cyp7b steroid hydroxylase-encoding DNA molecules comprising contiguous pairs of sequences selected from
- (i) the sequence designated "exon 3" in SEQ Id No 3,
  - (ii) the sequence designated "exon 4" in SEQ Id No 3, and
- (i) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecules defined in (h) under standard hybridization conditions defined as 2 x SSC at 65°C.
- (j) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecule defined in (h) or (i) under reduced stringency hybridization conditions defined as 6 x SSC at 55°C.

- (k) Coding sequences of DNA molecules comprising a contiguous coding sequence consisting of the sequences "exon 3" and "exon 4" in SEQ Id No 3, and
- 5 (l) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecules defined in (k) under standard hybridization conditions defined as 2 x SSC at 65°C.
- 10 (m) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecule defined in (k) or (l) under reduced stringency hybridization conditions defined as 6 x SSC at 55°C.

It will be appreciated that the DNA sequences referred to may consist of or be derived from genomic DNA, but typically would consist of or be derived from cDNA. Such sequences could be obtained by probing an appropriate library (cDNA or genomic) using hybridisation probes based upon the sequences provided according to the invention

15 of International patent application No PCT/GB95/02465, or they could be prepared by chemical synthesis or by ligation of sub-sequences.

In the above definitions, Cyp7b steroid hydroxylases have been defined in terms of DNA sequence information. The Cyp7b steroid hydroxylase enzyme used in accordance with the process of the invention may alternatively or additionally be defined by reference

20 to amino acid sequence information, e.g. the amino acid sequences contained in SEQ ID NO. 4, SEQ ID NO. 5 or SEQ ID NO 6.

Thus the Cyp7b steroid hydroxylase enzyme used in accordance with the process of the invention may have sequences matching one of said sequences exactly, or alternatively, the enzymes used may have sequences which differ from the aforementioned

25 sequences, provided that the capacity to catalyse the introduction of a 7 $\alpha$ -hydroxyl group is not eliminated.

Thus, for example, mutant enzymes may be produced by known methods, for example site-directed mutagenesis or other PCR-based procedures, and the expression

products tested for their capacity to catalyse the introduction of a 7 $\alpha$ -hydroxyl group in selected substrates in accordance with the procedures described herein.

Having regard to the degree of homology between the rat, mouse and human enzymes and known data relating to species divergence of hydroxylase enzymes, it is preferred that by comparison with the DNA sequences of SEQ ID NO. 1, SEQ ID NO. 2 and SEQ ID NO.3, the mutant enzymes should be encoded by sequences having at least 50% homology, more preferably at least 60% homology and most preferably at least 70% homology with said sequences over a length of 50 contiguous nucleotides.

Preferably the mutant enzymes are encoded by sequences having at least 60% homology with the entire coding sequence, more preferably at least 70%.

Alternatively, by comparison with the amino acid sequences of SEQ ID NO. 4, SEQ ID NO. 5 and SEQ ID NO.6, it is preferred that mutant enzymes should have at least 50% homology, more preferably at least 60% homology and most preferably at least 70% homology with said sequences over a length of 30 contiguous amino acids. Preferably the mutant enzymes have at least 60% homology and more preferably 70% homology or more with the entire amino acid sequence in each case.

It is however preferred that such mutant enzymes do not differ too drastically from the aforementioned sequences and in this regard, where amino acid substitutions are made, that the substituted amino acids are preferably so-called "synonymous" or "conservative" substitutions, i.e. hydrophilic, hydrophobic, basic and acidic amino acids should preferably be substituted by amino acids in the same class (see US 5380712).

More specifically, it is preferred that the mutant enzymes differ from the precise sequences of those described herein by not more than 20, preferably not more than 10 and most preferably not more than 5 amino acid substitutions, insertions or deletions.

The Cyp7b enzymes described herein may be used in toxicological and drug evaluation studies and such uses form further aspects of the invention. In a particularly preferred embodiment of this aspect of the invention, a cell line capable of expressing a Cyp7b enzyme is used as a basis of an assay for one or more Cyp7b substrates. Such cell lines have utility in toxicological and drug evaluation studies. Most preferably the cell line comprises a prokaryotic or eucaryotic cell line which has been transformed so as artificially to express a Cyp7b enzyme. Examples include bacteria, yeast and mammalian cells. Also

included are transgenic animals, at least one tissue of which (especially a non-brain tissue) expresses Cyp7b enzyme. Such transgenic animals may be produced by known methods for introducing foreign coding sequences into somatic or germ line cells.

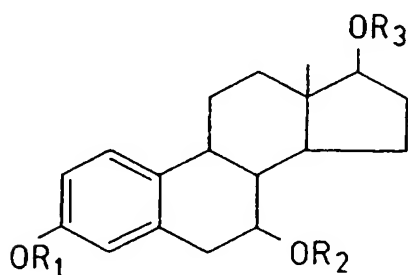
The substrates used in the method of the invention are characterised by possessing  
 5 a  $3\beta$ -hydroxyl group and further by preferably possessing the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol, with the proviso that where the substrate has the carbon skeleton of cholesterol, the substrate has a hydroxyl group in the 25, 26 or 27-position, preferably the 25-position.

Examples of such substrates include 25-hydroxycholesterol, dehydroepi-  
 10 androsterone, pregnenolone and estradiol, in which case the steroids produced will be  $7\alpha$ -hydroxy-25-hydroxycholesterol,  $7\alpha$ -hydroxydehydroepiandrosterone,  $7\alpha$ -hydroxy pregnenolone and  $7\alpha$ -hydroxyestradiol (i.e. estra 1,3,5(10)-triene-3,7 $\alpha$ ,17 $\beta$ -triol) respectively.

The  $7\alpha$ -hydroxylated steroid produced according to the invention may be oxidised  
 15 by known enzymatic or non-enzymatic procedures to produce 7-oxo substituted steroids and this further process step forms a further aspect of the invention.

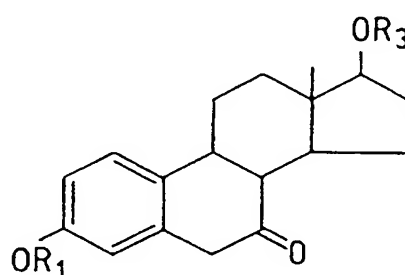
Certain  $7\alpha$ -hydroxy-substituted steroids produced according to the invention and certain corresponding 7-oxo derivatives are novel and provide a further aspect of the invention. Thus the present invention further provides novel  $3\beta$ -hydroxy steroids  
 20 characterised in that they have a  $7\alpha$ -hydroxy or 7-oxo substituent. Preferred novel steroids have the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol, with the proviso that where the skeleton is that of cholesterol, the 25, 26 or 27 position is hydroxylated, most preferably the 25 position.

Particular novel steroids are of the formula



25

Ia



Ib

wherein  $OR_1$ ,  $OR_2$  and  $OR_3$  each independently represents a free hydroxy group, an ether group or an esterified hydroxy group.

In the case where  $OR_1$ ,  $OR_2$  and  $OR_3$  each independently represents an ether group, each of  $R_1$ ,  $R_2$  and  $R_3$  may be selected from substituted or unsubstituted  $C_{1-6}$  alkyl groups, any such substituents being selected from OH, halogen (F, Cl, Br, I) amino,  $C_{1-6}$  alkylamino,  $C_{1-6}$  dialkylamino, COOH or  $COOR_4$  wherein  $R_4$  represents a  $C_{1-6}$  alkyl group which may be unsubstituted or substituted by one of the substituents referred to above.

In the case where  $OR_1$ ,  $OR_2$  and  $OR_3$  each independently represents an esterified hydroxy group, each of  $R_1$ ,  $R_2$  and  $R_3$  may have the formula  $R_5CO-$ , wherein  $R_5$  may be selected from substituted or unsubstituted  $C_{1-6}$  alkyl groups, any such substituents being selected from OH, halogen (F, Cl, Br, I) amino,  $C_{1-6}$  alkylamino,  $C_{1-6}$  dialkylamino, COOH or  $COOR_4$  wherein  $R_4$  represents a  $C_{1-6}$  alkyl group; and groups of the formula  $-OP(OH)_3$ . Where compounds of Formula Ia or Ib include substituents such as carboxyl groups, phosphate groups, or substituted or unsubstituted amino groups, the compounds may be in free form or in the form of acid addition salts with pharmacologically acceptable anions (such as, for example, phosphate or halide ions) or cations (such as, for example, alkaline metal cations). Thus, where  $OR_1$ ,  $OR_2$  or  $OR_3$  represents hemesuccinate  $HOOC(CH_2)_2CO$ , the resulting hemesuccinate may be in the form of, for example, an Na or K salt.

It will be realised that the present invention provides for  $7\alpha$ -hydroxylated and 7-oxo steroids as described above but which are further substituted at other positions directly on the steroid skeleton.

$7\alpha$ -Hydroxyestradiol and 7-oxoestradiol are specific examples of compounds of Formula Ia and Ib.

The invention will now be described in more detail with particular reference to the following Figures and Examples.

### Description of Figures

Figure 1 illustrates an autoradiogram of a TLC plate used in an experiment to assess the ability of various cell extracts to hydroxylate DHEA.

Figure 2 depicts the ability of various tissues to release radioactivity from 7-<sup>3</sup>H-pregnenolone.

Figure 3 illustrates the principal steroid interconversions mediated by Cyp7b.

Figure 4 is a histogram plotting fold induction of luciferase expression with  
5 concentration of various steroids as described in Example 5.

Figure 5 illustrates the attenuation of Cyp7b gene expression in Alzheimer's as described in Example 5.

Figure 6 shows mass spectrometer plots of 7 $\alpha$ -hydroxy-DHEA produced by the present process and a reference sample thereof.

## 10 **EXAMPLE 1 - Identification of substrate specificity of Mu Cyp7b**

### **A. Preparation of vaccinia expression construct**

To identify the reaction catalysed by Cyp7b a cDNA encoding the mouse enzyme, reported by Lathe, Rose and Stapleton (PCT/GB95/02465) and by Stapleton et al. (J. Biol. Chem. 270, 29739-1995, December 15 1995), was modified to introduce a translation  
15 initiation consensus sequence at the 5' end of the Cyp7b open reading frame as described therein. The modified cDNA was introduced into the genome of vaccinia virus by recombinational exchange according to standard procedures (see, for instance, Gonzalez et al., Meth. Enzymol. 206, 85-92, 1991 and references therein) as described in Lathe et al.

### 20 **B. Production of Cyp7b enzyme extracts.**

Hela cells were grown to semi confluence (10<sup>6</sup> cells per 5 cm dish; 5 ml medium) and infected with recombinant (VV-Cyp7b) and control (VV Copenhagen strain) vaccinia viruses at 0.1 pfu per cell; 16 hours later infected cells were washed and taken up into W (Waxman) buffer (0.1 M KP04, 1 mM EDTA, 20% glycerol pH 7.5; 500  $\mu$ l per plate) and  
25 recentrifuged (5 min., 1000 rpm).

For whole cell extracts cells were resuspended into 1/100 volume (50  $\mu$ l per plate) of W buffer and stored frozen at -70°C. For microsome preparation (Waxman, Biochem. J. 260, 81-85, 1989) cells were resuspended in 1/10 original volume of W buffer (500  $\mu$ l per plate); sonicated 6 x 5 seconds on ice, and unbroken cells were removed by  
30 centrifugation (10 min., 4°C, 3000 rpm).

The microsomal fraction was prepared from the supernatant by centrifugation

(100,000 g, 45 min., 4°C, Beckman SW50.1 rotor) and resuspended using a Potter homogeniser in 1/50 original volume of W buffer (100 µl per plate) before storage at -70°C.

- Control extracts were prepared from liver and brain from male rat by homogenising fresh tissue in W buffer (2.5 ml/g), clarifying briefly by centrifugation (4000 rpm, 5 min, 4°C); the supernatant was stored at -70°C.

### C. Substrate identification by thin-layer chromatography.

- $^{14}\text{C}$  or  $^3\text{H}$ -labelled steroids were purchased from DuPont-NEN ( $^{14}\text{C}$ -labelled molecules: specific activities 45-60 mCi/mmol.;  $^3\text{H}$ : specific activities 70-100 mCi/mmol). 1 nMol aliquots of labelled substrate were dried down, microsomes or cell and tissue extracts were added (25 to 50 µl), and diluted to a volume of 175 µl with W buffer.

- Reaction was started by the addition of 25 µl of 8 mM NADPH. After incubation at 37°C for 15 minutes the reaction was shaken with 500 µl of ethyl acetate (BDH). The organic phase was removed, dried down, and suspended into 10 µl ethyl acetate. Aliquots (5 x 2 µl) were applied to thin layer chromatography (TLC) sheets (Merck) and developed in ethyl acetate/n-hexane/acetic acid 16:8:1 (solvent system N of Waxman, Meth. Enzymol. 206, 462-476, 1991). After drying, chromatograms  $^{14}\text{C}$  were visualised by exposure to X-ray film.  $^3\text{H}$ -labelled chromatograms were treated with EN $^3$  HANCETM (DuPont-NEN) spray prior to exposure.

### D. Results

- Figure 1 is an autoradiogram of a TLC plate run in solvent system N; substrate was  $^3\text{H}$ -DHEA and samples were extracted with ethyl acetate and dried prior to application to the TLC plate (origin at bottom of figure). Extracts were 1, Microsomes from Hela cells infected with control vaccinia virus (negative control); 2, Microsomes from Hela cells infected with VVCyp7b; 3, Duplicate preparation of microsomes from Hela cells infected with VVCyp7b; 4, Rat brain homogenate.

- As can be seen from Figure 1, microsomes from cells infected with recombinant vaccinia expressing Cyp7b converted  $^{14}\text{C}$ -dehydroepiandrosterone (DHEA) to a lower mobility form most consistent with hydroxylation. Brain extracts yielded a product of indistinguishable mobility, consistent with our earlier demonstration that Cyp7b is



expressed in brain. From the relative mobility of the product we surmised that Cyp7b could be hydroxylating DHEA at the 7 position. Progesterone, corticosterone, cortisol and testosterone were at best inefficiently metabolised, if at all. However, pregnenolone and estradiol were both converted by the enzymes, as was 25-hydroxy cholesterol. All these  
5 substrates are distinguished by a 3 $\beta$  hydroxy group.

**EXAMPLE 2 - Identification of the position of the modification by  $^3\text{H}$ -release.**

To identify the position of the modification,  $^3\text{H}$ -pregnenolone (NEN) was employed in which the  $^3\text{H}$  substitution was predominantly at the 7 position on the steroid backbone. Microsomal extracts were incubated with  $^3\text{H}$ -pregnenolone under the same conditions as  
10 used earlier. Following reaction, labelled steroids were extracted with ethyl acetate (2 x 1 ml), and discarded; release of  $^3\text{H}$  into the aqueous phase was monitored by liquid scintillation counting.

Referring to Figure 2, 7- $^3\text{H}$ -pregnenolone was incubated with extracts and assayed for release of radioactivity into the aqueous phase following extraction with ethyl acetate.  
15 Extracts were 1, Microsomes from Hela cells infected with control vaccinia virus (negative control); 2, Microsomes from Hela cells infected with VVCyp7b; 3, Duplicate preparation of microsomes from Hela cells infected with VVCyp7b; 4, Rat brain homogenate; 5, Rat liver homogenate.

As seen in Figure 2 microsomes from cells infected with recombinant  
20 vaccinia expressing Cyp7b efficiently released  $^3\text{H}$  into the aqueous phase. Brain also performed this reaction but not liver. Release of  $^3\text{H}$  from the 7 position of pregnenolone demonstrates that Cyp7b hydroxylates pregnenolone at the 7-position to generate 7-hydroxy pregnenolone (7HP); it may be concluded that Cyp7b also hydroxylates DHEA (to generate 7-hydroxy DHEA [7HD]) and estradiol to generate 7-hydroxy estradiol [7HE].

25

**EXAMPLE 3 - Stereochemistry of the Cyp7b hydroxylation.**

Steroids hydroxylated at a variety of positions (egs. 2, 6, 7, 15, 16) differ in their mobility on TLC depending on whether the modification is in the  $\alpha$ - or the  $\beta$ -position (Waxman, Meth. Enzymol. 206, 462-476, 1991). Purified 7 $\alpha$ -hydroxy DHEA was  
30 obtained (kind gift of Dr. H.A. Lardy, Enzyme Institute, University of Wisconsin), mixed with the product of Cyp7b action on DHEA, and subjected to TLC. The product

comigrated with 7 $\alpha$ -hydroxy-DHEA, demonstrating that Cyp7b is a 7 $\alpha$  hydroxylase.

#### EXAMPLE 4 - Activity of enzyme in 7 $\alpha$ -hydroxylation of pregnenolone and DHEA

To examine the catalytic activity of the enzyme Cyp7b CDNAs were expressed in mammalian cell lines. Cell extracts showed substantial NADPH-dependent conversion of DHEA ( $K_m$  13.3 $\mu$ M;  $V_{max}$  288pmol/min/mg) and pregnenolone ( $K_m$  3.6 $\mu$ M;  $V_{max}$  34 pmol/min/mg) to slower migrating forms on thin layer chromatography. Products of identical mobility were generated by rat brain extracts. The expressed enzyme was less active against 25-hydroxycholesterol, 17 $\beta$ -estradiol and 5 $\alpha$ -androstane-3 $\beta$ , 17 $\beta$ -diol, with low to undetectable activity against progesterone, corticosterone and testosterone. When [3H-7 $\alpha$ ] pregnenolone was incubated with Cyp7b extracts the extent of release of radioactivity into the medium suggested that hydroxylation was preferentially at the 7 $\alpha$ -position. In gas chromatography and mass spectrometry of the modified steroid arising from incubation of DHEA with Cyp7b extracts, the retention time and fragmentation patterns were identical to those obtained with authentic 7 $\alpha$ -hydroxy DHEA (7HD); the reaction product also co-migrating with 7HD on TLC.

Mass spectrometry: A 10x scaled up reaction was employed using 95% unlabelled DHEA (Sigma) and 5% [14C]-DHEA (final specific activity 2.25-3mCi/mmol) and reaction time was extended to 1 hour. Product was purified by TLC, excised and extracted with ethyl acetate before drying down. The dried residue and authentic 7HD (50mg) were converted to their methoxime -trimethylsilyl (MO-TMS) derivatives. Analysis of these products was performed using a Trio 100 mass spectrometer operating in electron impact (EI) mode, linked to a HP5890 gas chromatograph fitted with a HP-1 cross-linked methyl siloxane column (25m, i.d. 0.25mm, 0.17 mm film) under the following conditions: electron energy 70eV, source temperature 200°C, interface temperature 280°C, oven temperature 50°C increasing at 30°C per minute to 200°C, and then 10°C per minute to 300°C, injection temperature 280°C.

#### EXAMPLE 5 - Cis-trans co-transfection assay; demonstration of antagonism.

Chinese hamster ovary (CHO) cells were maintained and transfected in Dulbecco's modification of Eagle's medium (DMEM) supplemented with 15% foetal bovine serum, 100IU/ml penicillin, 100 $\mu$ g/ml streptomycin and 200mM L-glutamine (all Gibco BRL,

Paisley, UK).

24 hours prior to transfection CHO cells were plated at a density of  $3 \times 10^5/60$  mm dish (Costar UK). Cells were transfected by the calcium phosphate method. Briefly, 5µg of MMTV-LUC and 1µg of pRShGR or 5µg of pSV2 as a control for transfection efficiency were made up to a total of 10µg/plate of DNA with pGEM3. 30µl of 2.5M CaCl<sub>2</sub> was diluted ten-fold with sterile water and 300µl was added to the DNA. Next 300µl of 2 x Hepes buffered saline (280 mM NaCl, 10mM KCl, 1.5mM Na<sub>2</sub>HPO<sub>4</sub>.2H<sub>2</sub>), 50mM Hepes, 12mM dextrose, pH 7.05) was added slowly with swirling to the DNA/CaCl<sub>2</sub> mixture. This solution was left for 30 minutes in order for a fine precipitate to form and 600µl was added dropwise to each plate. After 24 hours the medium was removed and the cells were washed in serum free medium and culture for a further 24 hours in medium containing 10% charcoal-stripped serum together with the appropriate concentrations of DHEA/7α-hydroxy-DHEA.

Six hours after the addition of DHEA/7α-hydroxyDHEA either B or Dex was added to each plate. The following day the cells were washed in PBS, lysed with 0.3ml of lysis buffer (25mM Tris-phosphate pH 7.8, 2mM DTT, 1% Triton X-100 and 10% glycerol), scraped, centrifuged and the supernatant assayed in duplicate in a Berthold luminometer in a total volume of 250µl, comprising 40µl of cell extract, 5µl of 30mM ATP, 100µl of assay buffer (20mM tricine, 1.07nM (MgCO<sub>3</sub>)<sub>4</sub>.Mg(OH)<sub>2</sub>.%H<sub>2</sub>O, 2.67mM MgSO<sub>4</sub>, 0.1mM EDTA, 33.3mM DTT, 0.2mg/ml coenzyme A) and 105µl luciferin (Promega UK) injected to initiate the reaction. Light emission was measured over 10 seconds and relative light units/microgram of protein was calculated.

Results are shown in Figure 4 wherein the fold induction of luciferase is illustrated by histogram for control, additions of DHEA, 7α-hydroxy-DHEA (7HD) alone and these additions in presence of an GR activating concentration of corticosterone. This result shows that 7HD, but not DHEA, acts as an antagonist of corticosterone effect in activating the GR-mediated transcription.

#### EXAMPLE 6 - Cyp7b expression in Alzheimers neurons

Cryostat brain sections (10µm) from control and Alzheimer's hippocampus were cut, thaw mounted onto gelatine-subbed poly-L-lysine coated slides and stored at -80°C.

For *in-situ* hybridization studies, brain sections were post-fixed in 4%

paraformaldehyde by acetylation (0.25% acetic anhydride in 0.1M triethanolamine, pH 8.0) for 10 minutes, rinsed in phosphate buffered saline, dehydrated through graded alcohols and air dried. Hybridization was carried out using 200µl of [<sup>35</sup>S]-UTP-labelled cRNA antisense probe (10 x 10<sup>6</sup> dpm/ml in hybridization buffer) synthesized *in vitro* from a 500  
5 bp XbaI/PstI fragment of the human Cyp7b pMMc1 clone linearised with XbaI and transcribed with T3 RNA for sense probes. Sections were prehybridized with 20µl of prehybridization buffer ( as hybridization buffer but omitting the dextran sulphate) per slide at 50°C for 3 hours.

Following hybridization with probe at 50°C overnight sections were treated with  
10 RNase A (30µg/ml, 45 minutes at 37°C) and washed to a final stringency of 0.1 x SSC at 60°C. Slides were dehydrated, dipped in photographic emulsion (NTB-2, Kodak) and exposed at 4°C for 5 weeks before being developed and counterstained with 1% pyronin. The density of silver grains was assessed over individual hippocampal neurons by computer-assisted grain counting using an image analysis system (Seescan plc, Cambridge,  
15 UK), with the analysis carried out blind (sections were cut and coded by a separate individual). For each slide, one hippocampal section represents each subject. 6-10 neurons/subregion were assessed and background, counted over areas of white matter, was subtracted. Data were assessed by ANOVA followed by Scheffe post hoc test. Significance was set at p<0.05. Values are means ± S.E.M.

20 Figure 5 is a histogram showing Cyp7b expression as indicated by grain count per neuron in the dentate gyrus, CA1 and CA3 subfields of Alzheimer's disease samples as compared to the age matched control brains.

## CONCLUSIONS

It can be concluded that Cyp7b, and cognate enzymes from rat, human and other  
25 mammalian species, are 7α-hydroxylases specific for steroid substrates with a 3β hydroxy group. While activities for 7-hydroxylating DHEA, pregnenolone and cholesterol have been recorded previously in a variety of crude tissue homogenates (eg. Akwa et al., Biochem. J. 288, 959-964, 1992) no characterisation of the enzyme responsible was performed previously and no activity on estradiol was recorded. Recombinant organisms expressing  
30 Cyp7b thus provide a route to the large scale manufacture of 7HP, 7HD, and 7HE, principally but not exclusively for therapeutic use or for the production of further steroid derivatives such as 7-oxo molecules.

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CLAIMS

1. The use of a  $7\alpha$ -hydroxy or 7-oxo substituted  $3\beta$ -hydroxy-steroid, or a derivative thereof substituted independently at one or both of the 7- and 3- positions with an ester or ether group, in the manufacture of a pharmaceutical composition for the therapy of neuropsychiatric, immune and/or endocrine disorders or for inducing cognitive enhancement.
2. The use according to Claim 1 wherein said disorders are selected from
- (a) deficits of cognition in aging
  - 10 (b) Alzheimer's disease
  - (c) deficits of immune system in aging
  - (d) deficits of immune function in HIV infection
  - (e) glucocorticoid or mineralocorticoid excess
  - (f) diabetes
  - 15 (g) depression
  - (h) osteoporosis and hypercalcemia
  - (I) hyperglycemia and hyperlipodemia
  - (j) muscle atrophy
  - (k) arterosclerosis
  - 20 (l) steroid diabetes
3. The use as claimed in claim 1 or claim 2 wherein the steroid has a  $3\beta$ -substituent-OR<sub>1</sub> and/or a  $7\alpha$ -substituent -OR<sub>2</sub> where -OR<sub>1</sub> and -OR<sub>2</sub> each independently represents a free hydroxy, ester or ether group,
- 25 wherein each of R<sub>1</sub> and R<sub>2</sub> are independently selected from the group consisting of hydrogen, substituted or unsubstituted C<sub>1-6</sub> alkyl groups, groups R<sub>5</sub>CO-, wherein R<sub>5</sub> may be selected from substituted or unsubstituted C<sub>1-6</sub> alkyl groups, and groups of the formula -OP(OH)<sub>3</sub>, wherein any substituents are selected from OH, halogen (F, Cl, Br, I) amino, C<sub>1-6</sub> alkylamino, C<sub>1-6</sub> dialkylamino, COOH or COOR<sub>4</sub> wherein R<sub>4</sub> represents a



C<sub>1-6</sub> alkyl group; and wherein the compounds may be in free form or in the form of acid addition salts with pharmacologically acceptable anions.

4. The use as claimed in any one of claims 1 to 3 characterised in that the steroid is one possessing the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol,
- 5 5. The use of a Cyp7b steroid hydroxylase enzyme in the manufacture of a test kit for use in the diagnosis of neuropsychiatric, immune and endocrine disorders.
6. The use according to Claim 5 wherein said disorders are selected from
  - (a) deficits of cognition in aging
  - (b) Alzheimer's disease
  - 10 (c) deficits of immune system in aging
  - (d) deficits of immune function in HIV infection
  - (e) glucocorticoid or mineralocorticoid excess
  - (f) diabetes
  - (g) depression
  - 15 (h) osteoporosis and hypercalcemia
  - (i) hyperglycemia and hyperlipodemia
  - (j) muscle atrophy
  - (k) arterosclerosis
  - (l) steroid diabetes
- 20 7. An antibody, especially a monoclonal antibody, characterised by specifically binding Cyp7b enzymes.
8. The use of an antibody as claimed in Claim 5 in a test kit for assaying for the presence of Cyp7b enzymes.
9. The use of Cyp7b coding sequences or antisense sequences in the manufacture of  
25 a targeted drug for gene therapy of Cyp deficiencies or excesses or for promoting immunogenic processes.

10. The use claimed in Claim 9 wherein a vector is co-administered with a vaccine formulation, whereby on administration, a Cyp7b sequence is expressed and the produced expression product augments an immunogenic property of the vaccine.
11. A process of producing a 7 $\alpha$ -hydroxy-substituted steroid which comprises  
5    subjecting a corresponding steroid substrate having no substituent in the 7-position to hydroxylation in the presence of a Cyp7b steroid hydroxylase enzyme.
12. A process according to Claim 11 wherein the enzyme is a mouse, rat or human Cyp7b steroid hydroxylase enzyme.
13. A process according to Claim 11 wherein the Cyp7b steroid hydroxylase enzyme  
10    has a sequence encoded by DNA coding sequences of Cyp7b enzymes selected from
- (a) Coding sequences of DNA molecules comprising the coding sequence for rat Cyp7b set forth in SEQ Id No: 1,
  - (b) Coding sequences of DNA molecules comprising the coding sequence for mouse Cyp7b set forth in SEQ Id No: 2,
  - 15    (c) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecule defined in (a) or (b) under standard hybridization conditions defined as 2 x SSC at 65°C.
  - (d) Cyp7b steroid hydroxylase-encoding DNA molecules capable of  
20    hybridizing with the DNA molecule defined in (a), (b) or (c) under reduced stringency hybridization conditions defined as 6 x SSC at 55°C.
14. A process according to Claim 11 wherein the Cyp7b steroid hydroxylase enzyme has a sequence encoded by DNA coding sequences of Cyp7b enzymes selected from

- (e) DNA coding sequences selected from the following:
- (i) the sequence designated "exon 3" in SEQ Id No 3.
  - (ii) the sequence designated "exon 4" in SEQ Id No 3. and
- 5 (f) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecules defined in (e) under standard hybridization conditions defined as 2 x SSC at 65°C.
- 10 (g) Cyp7b steroid hydroxylase encoding DNA molecules capable of hybridizing with the DNA molecule defined in (e) or (f) under reduced stringency hybridization conditions defined as 6 x SSC at 55°C.
- (h) Cyp7b steroid hydroxylase-encoding DNA molecules comprising contiguous pairs of sequences selected from
- (i) the sequence designated "exon 3" in SEQ Id No 3,
  - (ii) the sequence designated "exon 4" in SEQ Id No 3, and
- 15 (i) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecules defined in (h) under standard hybridization conditions defined as 2 x SSC at 65°C.
- 20 (j) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecule defined in (h) or (i) under reduced stringency hybridization conditions defined as 6 x SSC at 55°C.
- (k) Coding sequences of DNA molecules comprising a contiguous coding sequence consisting of the sequences "exon 3" and "exon 4" in SEQ Id No 3, and

- (l) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecules defined in (k) under standard hybridization conditions defined as 2 x SSC at 65°C.
- (m) Cyp7b steroid hydroxylase-encoding DNA molecules capable of hybridizing with the DNA molecule defined in (k) or (l) under reduced stringency hybridization conditions defined as 6 x SSC at 55°C.

15. A process according to Claim 11 wherein the Cyp7b steroid hydroxylase enzyme has a sequence encoded by DNA coding sequences of Cyp7b enzymes selected from the amino acid sequences contained in SEQ ID NO. 4, SEQ ID NO. 5 or SEQ ID NO. 6 or a sequence which has at least 50% homology with one or more of the aforementioned sequences, provided that the capacity to catalyse the introduction of a 7 $\alpha$ -hydroxyl group is not eliminated.

16. A process according to Claim 15 wherein the Cyp7b steroid hydroxylase enzyme has a sequence encoded by a DNA coding sequence which has at least 60% homology, and preferably at least 70% homology with one or more of the aforementioned sequences, provided that the capacity to catalyse the introduction of a 7 $\alpha$ -hydroxyl group is not eliminated.

17. A process according to Claim 15 wherein the Cyp7b steroid hydroxylase enzyme has a sequence which differs from the amino acid sequences contained in SEQ ID NO. 4, SEQ ID NO. 5 or SEQ ID NO. 6 by not more than 20, preferably not more than 10 and most preferably not more than 5 amino acid substitutions, insertions or deletions.

18. A process according to any preceding claim wherein substrate is a steroid possessing a 3 $\beta$ -hydroxyl group.

19. A process according to any preceding claim wherein the substrate is a steroid possessing the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol, with

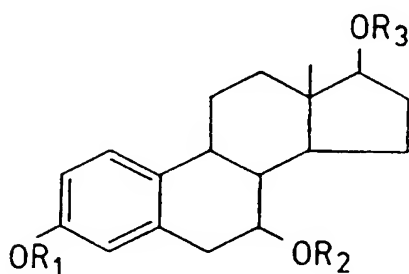
the proviso that where the substrate has the carbon skeleton of cholesterol, the substrate has a hydroxyl group in the 25, 26 or 27-position.

20. A process according to Claim 19 wherein the substrate is 25-hydroxycholesterol, dehydroepiandrosterone, pregnenolone or estradiol.

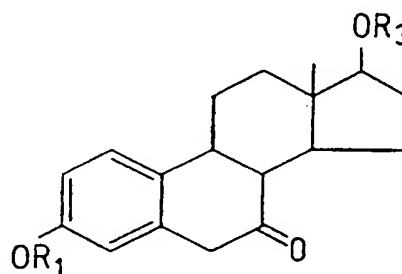
5 21. A process according to any preceding claim wherein the produced  $7\alpha$ -hydroxy-substituted steroid is  $7\alpha$ -hydroxyestradiol,  $7\alpha$ -hydroxypregnenolone or  $7\alpha$ -hydroxydehydroepiandrosterone.

22. A process according to any preceding claim wherein produced steroid is subjected to an oxidation step to convert an H.OH to an oxo group.

10 23. A steroid of the formula



Ia



Ib

wherein  $OR_1$ ,  $OR_2$  and  $OR_3$  each independently represents a free hydroxy group, an ether group or an esterified hydroxy group.

24. A steroid according to Claim 23 wherein  
 15 each of  $R_1$ ,  $R_2$  and  $R_3$  may be selected from substituted or unsubstituted  $C_{1-6}$  alkyl groups, any such substituents being selected from OH, halogen (F, Cl, Br, I) amino,  $(C_{1-6})$  alkylamino,  $C_{1-6}$  dialkylamino,  $COOH$  or  $COOR_4$  wherein  $R_4$  represents a  $C_{1-6}$  alkyl group which may be unsubstituted or substituted by one of the substituents referred to above; or

OR<sub>1</sub>, OR<sub>2</sub> and OR<sub>3</sub> each independently represents an esterified hydroxy group, of the formula R<sub>5</sub>COO-, wherein R<sub>5</sub> may be selected from substituted or unsubstituted C<sub>1-6</sub> alkyl groups, any such substituents being selected from OH, halogen (F, Cl, Br, I) amino, C<sub>1-6</sub> alkylamino, C<sub>1-6</sub> dialkylamino, COOH or COOR<sub>4</sub> wherein R<sub>4</sub> represents a C<sub>1-6</sub> alkyl group;  
5 or

OR<sub>1</sub>, OR<sub>2</sub> and OR<sub>3</sub> each independently represents an esterified hydroxy group of formula -OP(OH)<sub>3</sub>,  
or a pharmacologically acceptable salt of such a compound.

10 25. 7 $\alpha$ -Hydroxyestradiol or 7-oxoestradiol.

26. A steroid as claimed in Claim 23 characterised in that it is a 3 $\beta$ -hydroxy steroid.

27. A process of producing an oxo-substituted steroid which comprises subjecting 7 $\alpha$ -hydroxyestradiol, 7 $\alpha$ -hydroxypregnenolone or 7 $\alpha$ -hydroxydehydroepiandrosterone to oxidation.

15 28. A method for treating a human or animal requiring therapy for a neuropsychiatric, immune and endocrine disorder or for inducing cognitive enhancement comprising the administration of an effective amount of a 7 $\alpha$ -hydroxy or 7-oxo substituted 3 $\beta$ -hydroxy-steroid or derivative thereof independently substituted at one or both of the 7-and 3-positions by an ester or ether group.

20

29. A method according to Claim 28 wherein said disorders are selected from

- (a) deficits of cognition in aging
- (b) Alzheimer's disease
- (c) deficits of immune system in aging
- 25 (d) deficits of immune function in HIV infection
- (e) glucocorticoid or mineralocorticoid excess
- (f) diabetes
- (g) depression

- (h) osteoporosis and hypercalcemia
- (l) hyperglycemia and hyperlipodemia
- (j) muscle atrophy
- (k) arterosclerosis
- 5 (l) steroid diabetes

30. A method as claimed in claim 28 wherein the steroid possesses the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol and has a  $3\beta$ -substituent-OR<sub>1</sub> and/or a  $7\alpha$ -substituent -OR<sub>2</sub> where -OR<sub>1</sub> and -OR<sub>2</sub> each independently represents a free hydroxy, ester or ether group,

10 wherein each of R<sub>1</sub> and R<sub>2</sub> are independently selected from the group consisting of hydrogen, substituted or unsubstituted C<sub>1-6</sub> alkyl groups, groups R<sub>5</sub>CO-, wherein R<sub>5</sub> may be selected from substituted or unsubstituted C<sub>1-6</sub> alkyl groups, and groups of the formula -OP(OH)<sub>3</sub>, wherein any substituents are selected from OH, halogen (F, Cl, Br, I) amino, C<sub>1-6</sub> alkylamino, C<sub>1-6</sub> dialkylamino, COOH or COOR<sub>4</sub> wherein R<sub>4</sub> represents a C<sub>1-6</sub> alkyl group; and wherein the compounds may be in free form or in the form of acid addition salts with pharmacologically acceptable anions.

31. A  $7\alpha$ -hydroxy or 7-oxo substituted  $3\beta$ -hydroxy-steroid possessing the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol, or a derivative thereof substituted independently at one or both of the 7- and 3- positions with an ester or ether group for use in therapy.

32. A steroid as claimed in claim 31 selected from  $7\alpha$ -hydroxydehydroepiandrosterone,  $7\alpha$ -hydroxypregnenolone and  $7\alpha$ -hydroxycortisol.

25 33. A pharmaceutical composition characterised in that it comprises a  $7\alpha$ -hydroxy or 7-oxo substituted  $3\beta$ -hydroxy steroid possessing the carbon skeleton of cholesterol, androsterone, pregnenolone or estradiol, or a derivative thereof substituted independently at one or both of the 7- and 3- positions with an ester or ether group, in association with a pharmaceutically acceptable carrier or diluent in a sterile and pyrogen free form.

## SEQUENCE LISTING

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40 (C) CITY: Edinburgh  
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45 (ii) TITLE OF INVENTION: NEUROSTEROIDS



(iii) NUMBER OF SEQUENCES: 6

(iv) COMPUTER READABLE FORM:

- 5 (A) MEDIUM TYPE: Floppy disk  
 (B) COMPUTER: IBM PC compatible  
 (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)

(v) CURRENT APPLICATION DATA:

APPLICATION NUMBER: \*\*\*\*\*

10 (2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

- 15 (A) LENGTH: 1763 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: double  
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS  
 (B) LOCATION:1..1245

20 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

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GCC TTG GAG TAC CAG TAT GTA ATG AAA AAC CCA AAA CAA TTA AGC TTT
48
Ala Leu Glu Tyr Gln Tyr Val Met Lys Asn Pro Lys Gln Leu Ser Phe
  1             5             10             15

25 GAG AAG TTC AGC CGA AGA TTA TCA GCG AAA GCC TTC TCT GTC AAG AAG
96
Glu Lys Phe Ser Arg Arg Leu Ser Ala Lys Ala Phe Ser Val Lys Lys
    20             25             30

30 CTG CTA ACT AAT GAC GAC CTT AGC AAT GAC ATT CAC AGA GGC TAT CTT
Leu Leu Thr Asn Asp Asp Leu Ser Asn Asp Ile His Arg Gly Tyr Leu
    35             40             45
144

CTT TTA CAA GGC AAA TCT CTG GAT GGT CTT CTG GAA ACC ATG ATC CAA
Leu Leu Gln Gly Lys Ser Leu Asp Gly Leu Leu Glu Thr Met Ile Gln
    50             55             60
192

35 GAA GTA AAA GAA ATA TTT GAG TCC AGA CTG CTA AAA CTC ACA GAT TGG
Glu Val Lys Glu Ile Phe Glu Ser Arg Leu Leu Lys Leu Thr Asp Trp
    65             70             75             80
240

AAT ACA GCA AGA GTA TTT GAT TTC TGT AGT TCA CTG GTA TTT GAA ATC
Asn Thr Ala Arg Val Phe Asp Phe Cys Ser Ser Leu Val Phe Glu Ile
    85             90             95
288

40 ACA TTT ACA ACT ATA TAT GGA AAA ATT CTT GCT GCT AAC AAA AAA CAA
Thr Phe Thr Thr Ile Tyr Gly Lys Ile Leu Ala Ala Asn Lys Lys Gln
    336

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|    | 100   | 105 | 110 |      |
|----|---|-----|-----|------|
|    | ATT ATC AGT GAG CTG AGG GAT GAT TTT TTA AAA TTT GAT GAC CAT TTC |     |     | 384  |
|    | Ile Ile Ser Glu Leu Arg Asp Asp Phe Leu Lys Phe Asp Asp His Phe |     |     |      |
|    | 115   | 120 | 125 |      |
| 5  | CCA TAC TTA GTA TCT GAC ATA CCT ATT CAG CTT CTA AGA AAT GCA GAA |     |     | 432  |
|    | Pro Tyr Leu Val Ser Asp Ile Pro Ile Gln Leu Leu Arg Asn Ala Glu |     |     |      |
|    | 130   | 135 | 140 |      |
|    | TTT ATG CAG AAG AAA ATT ATA AAA TGT CTC ACA CCA GAA AAA GTA GCT |     |     | 480  |
|    | Phe Met Gln Lys Lys Ile Ile Lys Cys Leu Thr Pro Glu Lys Val Ala |     |     |      |
| 10 | 145   | 150 | 155 | 160  |
|    | CAG ATG CAA AGA CGG TCA GAA ATT GTT CAG GAG AGG CAG GAG ATG CTG |     |     | 528  |
|    | Gln Met Gln Arg Arg Ser Glu Ile Val Gln Glu Arg Gln Glu Met Leu |     |     |      |
|    | 165   | 170 | 175 |      |
|    | AAA AAA TAC TAC GGG CAT GAA GAG TTT GAA ATA GGA GCA CAT CAT CTT |     |     | 576  |
| 15 | Lys Lys Tyr Tyr Gly His Glu Glu Phe Glu Ile Gly Ala His His Leu |     |     |      |
|    | 180   | 185 | 190 |      |
|    | GGC TTG CTC TGG GCC TCT CTA GCA AAC ACC ATT CCA GCT ATG TTC TGG |     |     | 624  |
|    | Gly Leu Leu Trp Ala Ser Leu Ala Asn Thr Ile Pro Ala Met Phe Trp |     |     |      |
|    | 195   | 200 | 205 |      |
| 20 | GCA ATG TAT TAT CTT CTT CAG CAT CCA GAA GCT ATG GAA GTC CTG CGT |     |     | 672  |
|    | Ala Met Tyr Tyr Leu Leu Gln His Pro Glu Ala Met Glu Val Leu Arg |     |     |      |
|    | 210   | 215 | 220 |      |
|    | GAC GAA ATT GAC AGC TTC CTG CAG TCA ACA GGT CAA AAG AAA GGA CCT |     |     | 720  |
|    | Asp Glu Ile Asp Ser Phe Leu Gln Ser Thr Gly Gln Lys Lys Gly Pro |     |     |      |
| 25 | 225   | 230 | 235 | 240  |
|    | GGA ATT TCT GTC CAC TTC ACC AGA GAA CAA TTG GAC AGC TTG GTC TGC |     |     | 768  |
|    | Gly Ile Ser Val His Phe Thr Arg Glu Gln Leu Asp Ser Leu Val Cys |     |     |      |
|    | 245   | 250 | 255 |      |
|    | CTG GAA AGC GCT ATT CTT GAG GTT CTG AGG TTG TGC TCC TAC TCC AGC |     |     | 816  |
| 30 | Leu Glu Ser Ala Ile Leu Glu Val Leu Arg Leu Cys Ser Tyr Ser Ser |     |     |      |
|    | 260   | 265 | 270 |      |
|    | ATC ATC CGT GAA GTG CAA GAG GAT ATG GAT TTC AGC TCA GAG AGT AGG |     |     | 864  |
|    | Ile Ile Arg Glu Val Gln Glu Asp Met Asp Phe Ser Ser Glu Ser Arg |     |     |      |
|    | 275   | 280 | 285 |      |
| 35 | AGC TAC CGT CTG CGG AAA GGA GAC TTT GTA GCT GTC TTT CCT CCA ATG |     |     | 912  |
|    | Ser Tyr Arg Leu Arg Lys Gly Asp Phe Val Ala Val Phe Pro Pro Met |     |     |      |
|    | 290   | 295 | 300 |      |
|    | ATA CAC AAT GAC CCA GAA GTC TTC GAT GCT CCA AAG GAC TTT AGG TTT |     |     | 960  |
|    | Ile His Asn Asp Pro Glu Val Phe Asp Ala Pro Lys Asp Phe Arg Phe |     |     |      |
| 40 | 305   | 310 | 315 | 320  |
|    | GAT CGC TTC GTA GAA GAT GGT AAG AAG AAA ACA ACG TTT TTC AAA GGA |     |     | 1008 |
|    | Asp Arg Phe Val Glu Asp Gly Lys Lys Lys Thr Thr Phe Phe Lys Gly |     |     |      |

|    | 325  | 330 | 335 |      |
|----|--|-----|-----|------|
|    | GGA AAA AAG CTG AAG AGT TAC ATT ATA CCA TTT GGA CTT GGA ACA AGC    |     |     | 1056 |
|    | Gly Lys Lys Leu Lys Ser Tyr Ile Ile Pro Phe Gly Leu Gly Thr Ser    |     |     |      |
|    | 340  | 345 | 350 |      |
| 5  | AAA TGT CCA GGC AGA TAC TTT GCA ATT AAT GAA ATG AAG CTA CTA GTG    |     |     | 1104 |
|    | Lys Cys Pro Gly Arg Tyr Phe Ala Ile Asn Glu Met Lys Leu Leu Val    |     |     |      |
|    | 355  | 360 | 365 |      |
|    | ATT ATA CTT TTA ACT TAT TTT GAT TTA GAA GTC ATT GAC ACT AAG CCT    |     |     | 1152 |
|    | Ile Ile Leu Leu Thr Tyr Phe Asp Leu Glu Val Ile Asp Thr Lys Pro    |     |     |      |
| 10 | 370  | 375 | 380 |      |
|    | ATA GGA CTA AAC CAC AGT CGC ATG TTT CTG GGC ATT CAG CAT CCA GAC    |     |     | 1200 |
|    | Ile Gly Leu Asn His Ser Arg Met Phe Leu Gly Ile Gln His Pro Asp    |     |     |      |
|    | 385  | 390 | 395 | 400  |
|    | TCT GAC ATC TCA TTT AGG TAC AAG GCA AAA TCT TGG AGA TCC TGA        |     |     | 1245 |
| 15 | Ser Asp Ile Ser Phe Arg Tyr Lys Ala Lys Ser Trp Arg Ser *          |     |     |      |
|    | 405  | 410 | 415 |      |
|    | AAGGGTGGCA GAGAAGCTTA GCGGAATAAG GCTGCACATG CTGAGCTCTG TGATTTGCTG  |     |     | 1305 |
|    | TACTCCCCAA ATGCAGCCAC TATTCTTGTT TGTTAGAAAA TGGCAAATTT TTATTTGATT  |     |     | 1365 |
|    | GCGATCCATC CAGTTTGTTT TGGGTCACAA AACCTGTCAT AAAATAAAGC GCTGTCATGG  |     |     | 1425 |
| 20 | TGTAACAAAAA TGTCATGGCA ATCATTTCAG GATAAGGTAA AATAACGTTT TCAAGTTTGT |     |     | 1485 |
|    | ACTTACTATG ATTTTATCA TTTGTAGTGA ATGTGCTTTT CCAGTAATAA ATTTGCGCCA   |     |     | 1545 |
|    | GGGTGATTTT TTTTAATTAC TGAAATCCTC TAATATCGGT TTTATGTGCT GCCAGAAAAG  |     |     | 1605 |
|    | TGTGCCATCA ATGGACAGTA TAACAATTTT CAGTTTTCCA GAGAAGGGAG AAATTAAGCC  |     |     | 1665 |
|    | CCATGAGTTA CGCTGTATAA AATTGTTCTC TTCAACTATA ATATCAATAA TGTCTATATC  |     |     | 1725 |
| 25 | ACCAGGTTAC CTTTGCATTA AATCGAGTTT TGCAAAAG                          |     |     | 1763 |

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 1880 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 81..1604

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

GGCAGGCACA GCCTCTGGTC TAAGAAGAGA GGGCACTGTG CAGAAGCCAT CGCTCCCTAC  
60

AGAGCCGCCA GCTCGTCGGG ATG CAG GGA GCC ACG ACC CTA GAT GCC GCC 110  
Met Gln Gly Ala Thr Thr Leu Asp Ala Ala  
420 425

5 TCG CCA GGG CCT CTC GCC CTC CTA GGC CTT CTC TTT GCC GCC ACC TTA 158  
Ser Pro Gly Pro Leu Ala Leu Leu Gly Leu Leu Phe Ala Ala Thr Leu  
430 435 440

10 CTG CTC TCG GCC CTG TTC CTC CTC ACC CGG CGC ACC AGG CGC CCT CGT 206  
Leu Leu Ser Ala Leu Phe Leu Leu Thr Arg Arg Thr Arg Arg Pro Arg  
445 450 455

GAA CCA CCC TTG ATA AAA GGT TGG CTT CCT TAT CTT GGC ATG GCC CTG 254  
Glu Pro Pro Leu Ile Lys Gly Trp Leu Pro Tyr Leu Gly Met Ala Leu  
460 465 470

15 AAA TTC TTT AAG GAT CCG TTA ACT TTC TTG AAA ACT CTT CAA AGG CAA 302  
Lys Phe Phe Lys Asp Pro Leu Thr Phe Leu Lys Thr Leu Gln Arg Gln  
475 480 485

CAT GGT GAC ACT TTC ACT GTC TTC CTT GTG GGG AAG TAT ATA ACA TTT 350  
His Gly Asp Thr Phe Thr Val Phe Leu Val Gly Lys Tyr Ile Thr Phe  
490 495 500 505

20 GTT CTG AAC CCT TTC CAG TAC CAG TAT GTA ACG AAA AAC CCA AAA CAA 398  
Val Leu Asn Pro Phe Gln Tyr Gln Tyr Val Thr Lys Asn Pro Lys Gln  
510 515 520

TTA AGC TTT CAG AAG TTC AGC AGC CGA TTA TCA GCG AAA GCC TTC TCT 446  
Leu Ser Phe Gln Lys Phe Ser Ser Arg Leu Ser Ala Lys Ala Phe Ser  
525 530 535

25 GTA AAG AAG CTG CTT ACT GAT GAC GAC CTT AAT GAA GAC GTT CAC AGA 494  
Val Lys Lys Leu Leu Thr Asp Asp Asp Leu Asn Glu Asp Val His Arg  
540 545 550

GCC TAT CTA CTT CTA CAA GGC AAA CCT TTG GAT GCT CTT CTG GAA ACT 542  
Ala Tyr Leu Leu Leu Gln Gly Lys Pro Leu Asp Ala Leu Leu Glu Thr  
555 560 565

30 ATG ATC CAA GAA GTA AAA GAA TTA TTT GAG TCC CAA CTG CTA AAA ATC 590  
Met Ile Gln Glu Val Lys Glu Leu Phe Glu Ser Gln Leu Leu Lys Ile  
570 575 580 585

35 ACA GAT TGG AAC ACA GAA AGA ATA TTT GCA TTC TGT GGC TCA CTG GTA 638  
Thr Asp Trp Asn Thr Glu Arg Ile Phe Ala Phe Cys Gly Ser Leu Val  
590 595 600

TTT GAG ATC ACA TTT GCG ACT CTA TAT GGA AAA ATT CTT GCT GGT AAC 686  
Phe Glu Ile Thr Phe Ala Thr Leu Tyr Gly Lys Ile Leu Ala Gly Asn  
605 610 615

40 AAG AAA CAA ATT ATC AGT GAG CTA AGG GAT GAT TTT TTT AAA TTT GAT 734

|    |  |      |
|----|--|------|
|    | Lys Lys Gln Ile Ile Ser Glu Leu Arg Asp Asp Phe Phe Lys Phe Asp  |      |
|    | 620 625 630  |      |
| 5  | GAC ATG TTC CCA TAC TTA GTA TCT GAC ATA CCT ATT CAG CTT CTA AGA<br>Asp Met Phe Pro Tyr Leu Val Ser Asp Ile Pro Ile Gln Leu Leu Arg | 782  |
|    | 635 640 645  |      |
|    | AAT GAA GAA TCT ATG CAG AAG AAA ATT ATA AAA TGC CTC ACA TCA GAA  | 830  |
|    | Asn Glu Glu Ser Met Gln Lys Lys Ile Ile Lys Cys Leu Thr Ser Glu  |      |
|    | 650 655 660 665  |      |
| 10 | AAA GTA GCT CAG ATG CAA GGA CAG TCA AAA ATT GTT CAG GAA AGC CAA  | 878  |
|    | Lys Val Ala Gln Met Gln Gly Gln Ser Lys Ile Val Gln Glu Ser Gln  |      |
|    | 670 675 680  |      |
|    | GAT CTG CTG AAA AGA TAC TAT AGG CAT GAC GAT TCT GAA ATA GGA GCA  | 926  |
|    | Asp Leu Leu Lys Arg Tyr Tyr Arg His Asp Asp Ser Glu Ile Gly Ala  |      |
|    | 685 690 695  |      |
| 15 | CAT CAT CTT GGC TTT CTC TGG GCC TCT CTA GCA AAC ACC ATT CCA GCT  | 974  |
|    | His His Leu Gly Phe Leu Trp Ala Ser Leu Ala Asn Thr Ile Pro Ala  |      |
|    | 700 705 710  |      |
| 20 | ATG TTC TGG GCA ATG TAT TAT ATT CTT CGG CAT CCT GAA GCT ATG GAA  | 1022 |
|    | Met Phe Trp Ala Met Tyr Tyr Ile Leu Arg His Pro Glu Ala Met Glu  |      |
|    | 715 720 725  |      |
|    | GCC CTG CGT GAC GAA ATT GAC AGT TTC CTG CAG TCA ACA GGT CAA AAG  | 1070 |
|    | Ala Leu Arg Asp Glu Ile Asp Ser Phe Leu Gln Ser Thr Gly Gln Lys  |      |
|    | 730 735 740 745  |      |
| 25 | AAA GGG CCT GGA ATT TCA GTC CAC TTC ACC AGA GAA CAA TTG GAC AGC  | 1118 |
|    | Lys Gly Pro Gly Ile Ser Val His Phe Thr Arg Glu Gln Leu Asp Ser  |      |
|    | 750 755 760  |      |
|    | TTG GTC TGC CTG GAA AGC ACT ATT CTT GAG GTT CTG AGG CTG TGC TCA  | 1166 |
|    | Leu Val Cys Leu Glu Ser Thr Ile Leu Glu Val Leu Arg Leu Cys Ser  |      |
|    | 765 770 775  |      |
| 30 | TAC TCC AGC ATC ATC CGA GAA GTG CAG GAG GAT ATG AAT CTC AGC TTA  | 1214 |
|    | Tyr Ser Ser Ile Ile Arg Glu Val Gln Glu Asp Met Asn Leu Ser Leu  |      |
|    | 780 785 790  |      |
| 35 | GAG AGT AAG AGT TTC TCT CTG CGG AAA GGA GAT TTT GTA GCC CTC TTT  | 1262 |
|    | Glu Ser Lys Ser Phe Ser Leu Arg Lys Gly Asp Phe Val Ala Leu Phe  |      |
|    | 795 800 805  |      |
|    | CCT CCA CTC ATA CAC AAT GAC CCG GAA ATC TTC GAT GCT CCA AAG GAA  | 1310 |
|    | Pro Pro Leu Ile His Asn Asp Pro Glu Ile Phe Asp Ala Pro Lys Glu  |      |
|    | 810 815 820 825  |      |
| 40 | TTT AGG TTC GAT CGG TTC ATA GAA GAT GGT AAG AAG AAA AGC ACG TTT  | 1358 |
|    | Phe Arg Phe Asp Arg Phe Ile Glu Asp Gly Lys Lys Lys Ser Thr Phe  |      |
|    | 830 835 840  |      |
|    | TTC AAA GGA GGG AAG AGG CTG AAG ACT TAC GTT ATG CCT TTT GGA CTC  | 1406 |
|    | Phe Lys Gly Gly Lys Arg Leu Lys Thr Tyr Val Met Pro Phe Gly Leu  |      |

|    | 845   | 850 | 855 |      |
|----|---|-----|-----|------|
|    | GGA ACA AGC AAA TGT CCA GGG AGA TAT TTT GCA GTG AAC GAA ATG AAG   |     |     | 1454 |
|    | Gly Thr Ser Lys Cys Pro Gly Arg Tyr Phe Ala Val Asn Glu Met Lys   |     |     |      |
|    | 860   | 865 | 870 |      |
| 5  | CTA CTG CTG ATT GAG CTT TTA ACT TAT TTT GAT TTA GAA ATT ATC GAC   |     |     | 1502 |
|    | Leu Leu Leu Ile Glu Leu Leu Thr Tyr Phe Asp Leu Glu Ile Ile Asp   |     |     |      |
|    | 875   | 880 | 885 |      |
|    | AGG AAG CCT ATA GGG CTA AAT CAC AGT CGG ATG TTT TTA GGT ATT CAG   |     |     | 1550 |
|    | Arg Lys Pro Ile Gly Leu Asn His Ser Arg Met Phe Leu Gly Ile Gln   |     |     |      |
| 10 | 890   | 895 | 900 | 905  |
|    | CAC CCC GAT TCT GCC GTC TCC TTT AGG TAC AAA GCA AAA TCT TGG AGA   |     |     | 1598 |
|    | His Pro Asp Ser Ala Val Ser Phe Arg Tyr Lys Ala Lys Ser Trp Arg   |     |     |      |
|    | 910   | 915 | 920 |      |
|    | -----   |     |     |      |
|    | AGC TGA AAGTGTGGCA GAGAAGCTTT GCAGAGTAAG GCTGCATGTG CTGAGCTCCG    |     |     | 1654 |
| 15 | Ser *   |     |     |      |
|    | TGATTTGGTG CACTCCCCCA AATGCAACCG CTACTCTTGT TTGAAAATGG CAAATTTATA |     |     | 1714 |
|    | TTTGGTTGAG ATCAATCCAG TTGGTTTTGG GTCACAAAAC CTGTCATAAA ATAAAGCAGT |     |     | 1774 |
|    | GTGATGGTTT AAAAAATGTC ATGGCAATCA TTTCAGGATA AGGTAAAATA ACATTTTCAA |     |     | 1834 |
| 20 | GTTTGTACTT ACTATGATTT TTATCATTTG TAGTGAATGT GCTTTT                |     |     | 1880 |

## (2) INFORMATION FOR SEQ ID NO: 3:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 3846 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: double
  - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

- (ix) FEATURE:
- (A) NAME/KEY: CDS
  - (B) LOCATION: 831..2078

- (ix) FEATURE:
- (A) NAME/KEY: exon (3)
  - (B) LOCATION: 831..1422

- (ix) FEATURE:
- (A) NAME/KEY: intron
  - (B) LOCATION: 1423..1872

## (ix) FEATURE:

(A) NAME/KEY: exon (4)

(B) LOCATION: 1873..2078

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

5 GGATCCAACC AAGTTTCCAG ATCTTATAAA TGTGGTGAAT GGTGAATGAC TTCCTGAAGA  
60

ATGGATGAAT GGATGTGTTT TAGTTTGGAA TCCTGTGTCA GTCACAAGTC AATATGTGAC 120

CTTGAACATG TTATTAAATC TCCCACATCC ATAAAAGTGA AAATGCTGGC ATTAGTGGAT 180

TTTTGCCAGT GTTGAATTAG ACATTTATTT GTGAGTACCT GCTCCATACA GTATGGTCAT 240

10 TTATTTGAGT TAAAATTGTT GTATTTGAAC AAAACTCAGA TGACACCTAA GCATGAAAAA 300

GCTCTTTATG AAGTATAAAT ACTCAGAAAT GGAATGGCAT GTTGCCAATT TGTCTTCTGC 360

TTTATTGAGG GAAATATATG AGAAGTATTT AAGTCAGGGG ATTATGAGGA ATATTTAAAG 420

GATANNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN 480

NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN 540

15 NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN 600

NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNTCTAGA GTGTTTTCCA CCATCTTTCA 660

AAGGAAACAT GTAGTGTAAC TTCGAATGAA ATGGATTTGT ATTAACTTT TTGCCTTAGT 720

TATTAGGGTC TTTCTAATTT TTGATTAACA TATTTTTTTA ATTTGTGGTG TTTATTTCTG 780

TTTTTATTAA CAAACGAACT CATATGCTCC TCTCTCTTTT TTTTTTTTCT GGAAAGTACA 840

20 TAACATTTAT ACCTGGACCC TTCCAGTACC AGCTAGTGAT AAAAAATCAT AAACAATTAA 900

GCTTTGAGT ATCTTCTAAT AAATTATCAG AGAAAGCATT TAGCATCAGT CAGTTGCAAA 960

AAAATCATGA CATGAATGAT GAGCTTCACC TCTGCTATCA ATTTTGTCAA GGCAAATCTT 1020

TGGACATACT CTTGGAAAGC ATGATGCAGA ATCTAAAACA AGTTTTTGAA CCCCAGCTGT 1080

TAAAAACCAC AAGTTGGGAC ACGGCAGAAC TGTATCCATT CTGCAGCTCA ATAATATTTG 1140

25 AGATCACATT TACAACTATA TATGGAAAAG TTATTGTTTG TGACAACAAC AAATTTATTA 1200

GTGAGCTAAG AGATGATTTT TAAAATTTG ATGACAAGTT TGCATATTTA GATCCAACA 1260

TACCCATTGA GCTTCTAGGA AATGTCAAGT CTATTAGAGA GAAAATTATA AAATGCTTCT 1320

CATCAGAAAA GTTAGCCAAG ATGCAAGGAT GGTGAGAAGT TTTTCAAAGC AGGCAAGATG 1380

ACCTGGAGAA ATATTATGTG CACGAGGACC TTGAAATAGG AGGTAAGAAC TTCTGAATGA 1440

30 GCACTTGCCT AAATAAAAAAT CATTTACATA GACCTCTGAA ATAAAAAAG ACAAATGGC 1500

|    |  |      |
|----|--|------|
| 7  | GACCTTGAAA ATTTTTTTAT GCTCTTTCTA ATTGGCTAAT GATAAATGTT TACTCTGATA  | 1560 |
|    | TAACCTCTAT AATTGATATT TTTTTTTTGT CTGAGGTGGT AAACAGATAC TTAATGGTGA  | 1620 |
|    | TAATGAGAAA GCGTATAACT AAGCTGCATT TATCCCTCTT ATCTCATCCC CGACCACACC  | 1680 |
| 5  | GCCCCCCCCA TACACATTAC ATTTTAAACT ATTCTCATTAG AGCAGAAAAT TAGACTTCAG | 1740 |
|    | AAGCCTATTG GTTCTCATTG GCATGCAGTG ATCCTTGGCT GGTCTGTGTC CTAACATCTT  | 1800 |
|    | TTAATTAGCA CACTGCAAAT CTAATCAGTG TAATAAACGC TATTAATCTT CCTTTTACACT | 1860 |
|    | TATTTTCTCC CACACATCAT TTAGGCTTTC TCTGGGCCTC TGTGGCAAAC ACTATTCCAA  | 1920 |
|    | CTATGTTCTG GGCAACGTAT TATCTTCTGC GGCACCCAGA AGCTATGGCA GCAGTGCGTG  | 1980 |
| 10 | ACGAAATTGA CCGTTTGCTG CAGTCAACAG GTCAAAGGA AGGGTCTGGA TTTCCCATCC   | 2040 |
|    | ACCTCACCAG AGAACAATTG GACAGCCTAA TCTGCCTAGG TAATTATTTT ATCTGTTATG  | 2100 |
|    | AAGAAAGAAG GTACCTCTCT GCAAACCTCGG TTTATCACTC ATAGCTGTTT ACAAGAGGTA | 2160 |
|    | GAGGACACAG CTGCTAATTG ACATAATAAC TCCCATTTAC ATCAATTATA AATTATGTAG  | 2220 |
|    | TTTATAGCCG TAGATCATCT CATTGCATGT AAACATAAGG CCTATGTAAT TAACTGTGTA  | 2280 |
| 15 | ATGTATGTAA AATTCTAACC AAAGCTTNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2340 |
|    | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2400 |
|    | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2460 |
|    | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2520 |
|    | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2580 |
|    | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2640 |
| 20 | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2700 |
|    | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2760 |
|    | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2820 |
|    | NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN NNNNNNNNNN  | 2880 |
| 25 | AAGTTAAATT CCATACCAAT GAGTTATTCT CT CTC TGTATTGACA TTTCATCTGC      | 2940 |
|    | GGTATCCTTT AGGGTACAAT GAGTTATTCT CTA CTC TGTATTGACA TTTCATCTGC     | 3000 |
|    | GGTATCCTTT AGGGTACAAT ATTCCAAGTT TCTTTAGACA AACGCAGGAA CAAATGTTCA  | 3060 |
|    | CATATTTCTG TTTCTTTATT CCTTTGACAA GTAGGCGAGC ATTTTAGCCT ATGTTGGTCT  | 3120 |
|    | CAAAAAAAT CTTTAAATA TGTTCCAGGT TCTTTAATGG GACCTTTCAG GAGCAAAAGT    | 3180 |



CCTCCCAGGT TTGGTCAATG TTCACCCTCN GTGGCCATTG AGGAAAATGC CCNNNNNGTT 3240  
 CTAGAGATTG TTCTCACTTC TCAGGCTAAG GCCCATTGAG CAATGCCAGA AAGCATGCCT 3300  
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 AAAGTAGTTG TAATTCATCC ACTCTTTTTA CTTTCAACTT TTTGCTATTA AAAAATCATT 3720  
 10 TTTAAATTTT AGTATTAAAG CAGAAACATT TAAATTTATT AGACCAGAAA AATAACAGAT 3780  
 TCTAGAACTA TAATTTGAAT CCATTTAAGC CCATAGCTAG AGCTAGAGAT TTTCACTATT 3840  
 GGATCC 3846

## (2) INFORMATION FOR SEQ ID NO: 4:

15 (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 415 amino acids  
 (B) TYPE: amino acid  
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

20 Ala Leu Glu Tyr Gln Tyr Val Met Lys Asn Pro Lys Gln Leu Ser Phe  
 1 5 10 15  
 Glu Lys Phe Ser Arg Arg Leu Ser Ala Lys Ala Phe Ser Val Lys Lys  
 20 25 30  
 25 Leu Leu Thr Asn Asp Asp Leu Ser Asn Asp Ile His Arg Gly Tyr Leu  
 35 40 45  
 Leu Leu Gln Gly Lys Ser Leu Asp Gly Leu Leu Glu Thr Met Ile Gln  
 50 55 60  
 Glu Val Lys Glu Ile Phe Glu Ser Arg Leu Leu Lys Leu Thr Asp Trp  
 65 70 75 80  
 30 Asn Thr Ala Arg Val Phe Asp Phe Cys Ser Ser Leu Val Phe Glu Ile  
 85 90 95  
 Thr Phe Thr Thr Ile Tyr Gly Lys Ile Leu Ala Ala Asn Lys Lys Gln  
 100 105 110  
 Ile Ile Ser Glu Leu Arg Asp Asp Phe Leu Lys Phe Asp Asp His Phe

|    | 115   | 120                             | 125                 |
|----|---|---------------------------------|---------------------|
|    | Pro Tyr Leu Val Ser Asp Ile                                     | Pro Ile Gln Leu                 | Leu Arg Asn Ala Glu |
|    | 130   | 135                             | 140                 |
| 5  | Phe Met Gln Lys Lys Ile Ile                                     | Lys Cys Leu Thr                 | Pro Glu Lys Val Ala |
|    | 145   | 150                             | 155 160             |
|    | Gln Met Gln Arg Arg Ser Glu Ile Val                             | Gln Glu Arg Gln Glu Met Leu     |                     |
|    |   | 165 170                         | 175                 |
|    | Lys Lys Tyr Tyr Gly His Glu Glu                                 | Phe Glu Ile Gly Ala His His Leu |                     |
|    |   | 180 185                         | 190                 |
| 10 | Gly Leu Leu Trp Ala Ser Leu Ala Asn Thr Ile                     | Pro Ala Met Phe Trp             |                     |
|    |   | 195 200                         | 205                 |
|    | Ala Met Tyr Tyr Leu Leu Gln His Pro Glu Ala                     | Met Glu Val Leu Arg             |                     |
|    |   | 210 215                         | 220                 |
| 15 | Asp Glu Ile Asp Ser Phe Leu Gln Ser Thr                         | Gly Gln Lys Lys Gly Pro         |                     |
|    |   | 225 230 235                     | 240                 |
|    | Gly Ile Ser Val His Phe Thr Arg Glu Gln Leu Asp Ser Leu Val Cys |                                 |                     |
|    |   | 245 250                         | 255                 |
|    | Leu Glu Ser Ala Ile Leu Glu Val Leu Arg Leu Cys Ser Tyr Ser Ser |                                 |                     |
|    |   | 260 265                         | 270                 |
| 20 | Ile Ile Arg Glu Val Gln Glu Asp Met Asp Phe Ser Ser Glu Ser Arg |                                 |                     |
|    |   | 275 280                         | 285                 |
|    | Ser Tyr Arg Leu Arg Lys Gly Asp Phe Val Ala Val Phe Pro Pro Met |                                 |                     |
|    |   | 290 295                         | 300                 |
| 25 | Ile His Asn Asp Pro Glu Val Phe Asp Ala Pro Lys Asp Phe Arg Phe |                                 |                     |
|    |   | 305 310 315                     | 320                 |
|    | Asp Arg Phe Val Glu Asp Gly Lys Lys Lys Thr Thr Phe Phe Lys Gly |                                 |                     |
|    |   | 325 330                         | 335                 |
|    | Gly Lys Lys Leu Lys Ser Tyr Ile Ile Pro Phe Gly Leu Gly Thr Ser |                                 |                     |
|    |   | 340 345                         | 350                 |
| 30 | Lys Cys Pro Gly Arg Tyr Phe Ala Ile Asn Glu Met Lys Leu Leu Val |                                 |                     |
|    |   | 355 360                         | 365                 |
|    | Ile Ile Leu Leu Thr Tyr Phe Asp Leu Glu Val Ile Asp Thr Lys Pro |                                 |                     |
|    |   | 370 375                         | 380                 |
| 35 | Ile Gly Leu Asn His Ser Arg Met Phe Leu Gly Ile Gln His Pro Asp |                                 |                     |
|    |   | 385 390 395                     | 400                 |
|    | Ser Asp Ile Ser Phe Arg Tyr Lys Ala Lys Ser Trp Arg Ser *       |                                 |                     |
|    |   | 405 410                         | 415                 |

## (2) INFORMATION FOR SEQ ID NO: 5:

## (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 508 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

Met Gln Gly Ala Thr Thr Leu Asp Ala Ala Ser Pro Gly Pro Leu Ala  
 1 5 10 15  
 10 Leu Leu Gly Leu Leu Phe Ala Ala Thr Leu Leu Leu Ser Ala Leu Phe  
 20 25 30  
 Leu Leu Thr Arg Arg Thr Arg Arg Pro Arg Glu Pro Pro Leu Ile Lys  
 35 40 45  
 15 Gly Trp Leu Pro Tyr Leu Gly Met Ala Leu Lys Phe Phe Lys Asp Pro  
 50 55 60  
 Leu Thr Phe Leu Lys Thr Leu Gln Arg Gln His Gly Asp Thr Phe Thr  
 65 70 75 80  
 Val Phe Leu Val Gly Lys Tyr Ile Thr Phe Val Leu Asn Pro Phe Gln  
 85 90 95  
 20 Tyr Gln Tyr Val Thr Lys Asn Pro Lys Gln Leu Ser Phe Gln Lys Phe  
 100 105 110  
 Ser Ser Arg Leu Ser Ala Lys Ala Phe Ser Val Lys Lys Leu Leu Thr  
 115 120 125  
 25 Asp Asp Asp Leu Asn Glu Asp Val His Arg Ala Tyr Leu Leu Leu Gln  
 130 135 140  
 Gly Lys Pro Leu Asp Ala Leu Leu Glu Thr Met Ile Gln Glu Val Lys  
 145 150 155 160  
 Glu Leu Phe Glu Ser Gln Leu Leu Lys Ile Thr Asp Trp Asn Thr Glu  
 165 170 175  
 30 Arg Ile Phe Ala Phe Cys Gly Ser Leu Val Phe Glu Ile Thr Phe Ala  
 180 185 190  
 Thr Leu Tyr Gly Lys Ile Leu Ala Gly Asn Lys Lys Gln Ile Ile Ser  
 195 200 205  
 35 Glu Leu Arg Asp Asp Phe Phe Lys Phe Asp Asp Met Phe Pro Tyr Leu  
 210 215 220  
 Val Ser Asp Ile Pro Ile Gln Leu Leu Arg Asn Glu Glu Ser Met Gln  
 225 230 235 240  
 Lys Lys Ile Ile Lys Cys Leu Thr Ser Glu Lys Val Ala Gln Met Gln

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## (2) INFORMATION FOR SEQ ID NO: 6:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 266 amino acids  
 (B) TYPE: amino acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: peptide

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

10 Gly Lys Tyr Ile Thr Phe Ile Pro Gly Pro Phe Gln Tyr Gln Leu Val  
 1 5 10 15  
 Ile Lys Asn His Lys Asn Leu Ser Phe Arg Val Ser Ser Asn Lys Leu  
 20 25 30  
 Ser Glu Lys Ala Phe Ser Ile Ser Gln Leu Gln Lys Asn His Asp Met  
 35 40 45  
 15 Asn Asp Glu Leu His Leu Cys Tyr Gln Phe Leu Gln Gly Lys Ser Leu  
 50 55 60  
 Asp Ile Leu Leu Glu Ser Met Met Gln Asn Leu Lys Gln Val Phe Glu  
 65 70 75 80  
 20 Pro Gln Leu Leu Lys Thr Thr Ser Trp Asp Thr Ala Glu Leu Tyr Pro  
 85 90 95  
 Phe Cys Ser Ser Ile Ile Phe Glu Ile Thr Phe Thr Thr Ile Tyr Gly  
 100 105 110  
 Lys Val Ile Val Cys Asp Asn Asn Lys Phe Ile Ser Glu Leu Arg Asp  
 115 120 125  
 25 Asp Phe Leu Lys Phe Asp Asp Lys Phe Ala Tyr Leu Val Ser Asn Ile  
 130 135 140  
 Pro Ile Glu Leu Leu Gly Asn Val Lys Ser Ile Arg Glu Lys Ile Ile  
 145 150 155 160  
 30 Lys Cys Phe Ser Ser Glu Lys Leu Ala Lys Met Gln Gly Trp Ser Glu  
 165 170 175  
 Val Phe Gln Ser Arg Gln Asp Asp Leu Glu Lys Tyr Tyr Val His Glu  
 180 185 190  
 Asp Leu Glu Ile Gly Ala His His Phe Gly Phe Leu Trp Val Ser Val  
 195 200 205  
 35 Ala Ser Thr Ile Pro Thr Met Phe Trp Ala Thr Tyr Tyr Leu Leu Arg  
 210 215 220

His Pro Glu Ala Met Ala Ala Val Arg Asp Glu Ile Asp Arg Leu Leu  
225 230 235 240

Gln Ser Thr Gly Gln Lys Glu Gly Ser Gly Phe Pro Ile His Leu Thr  
245 250 255

5 Arg Glu Gln Leu Asp Ser Leu Ile Cys Leu  
260 265

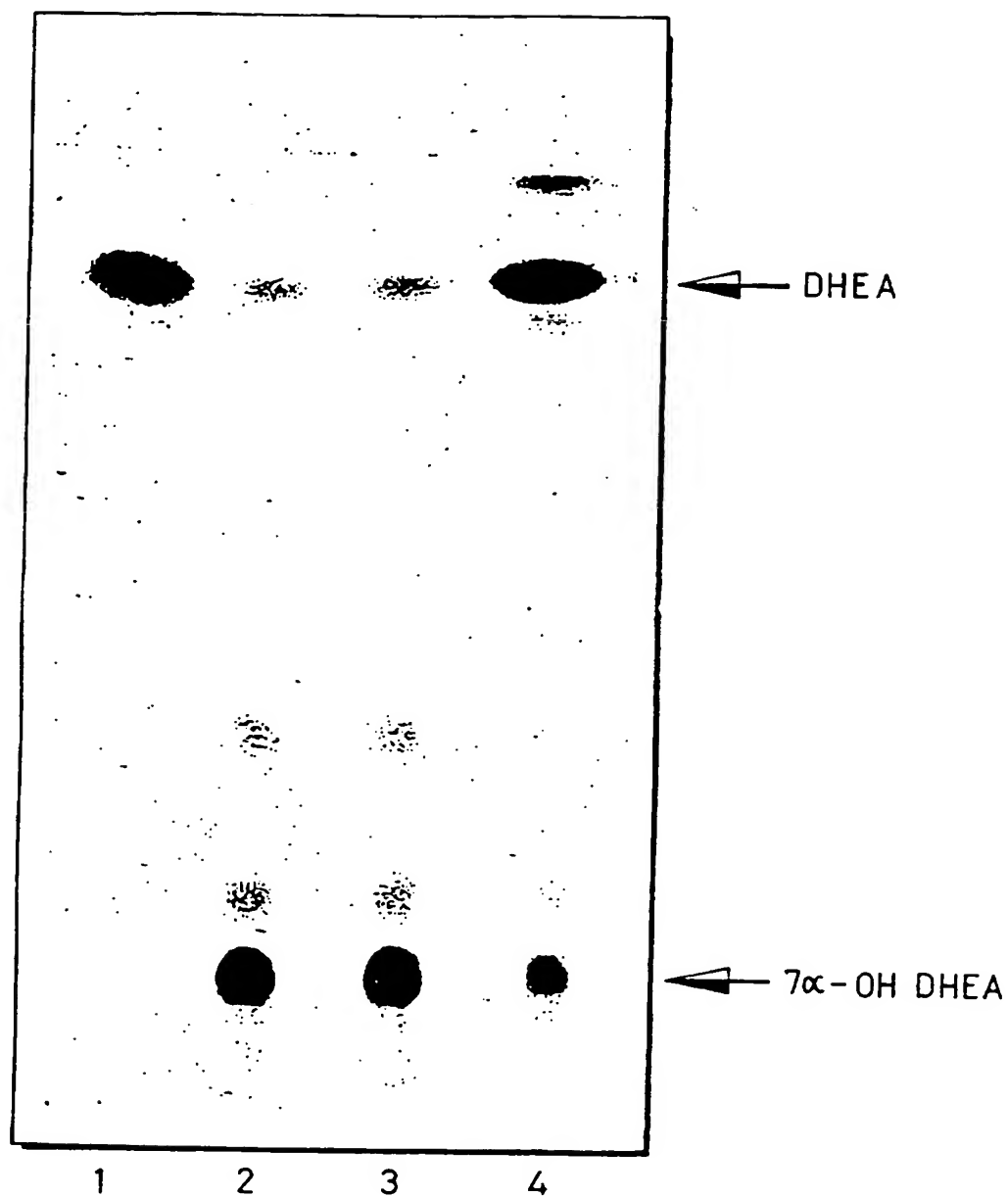


Fig. 1

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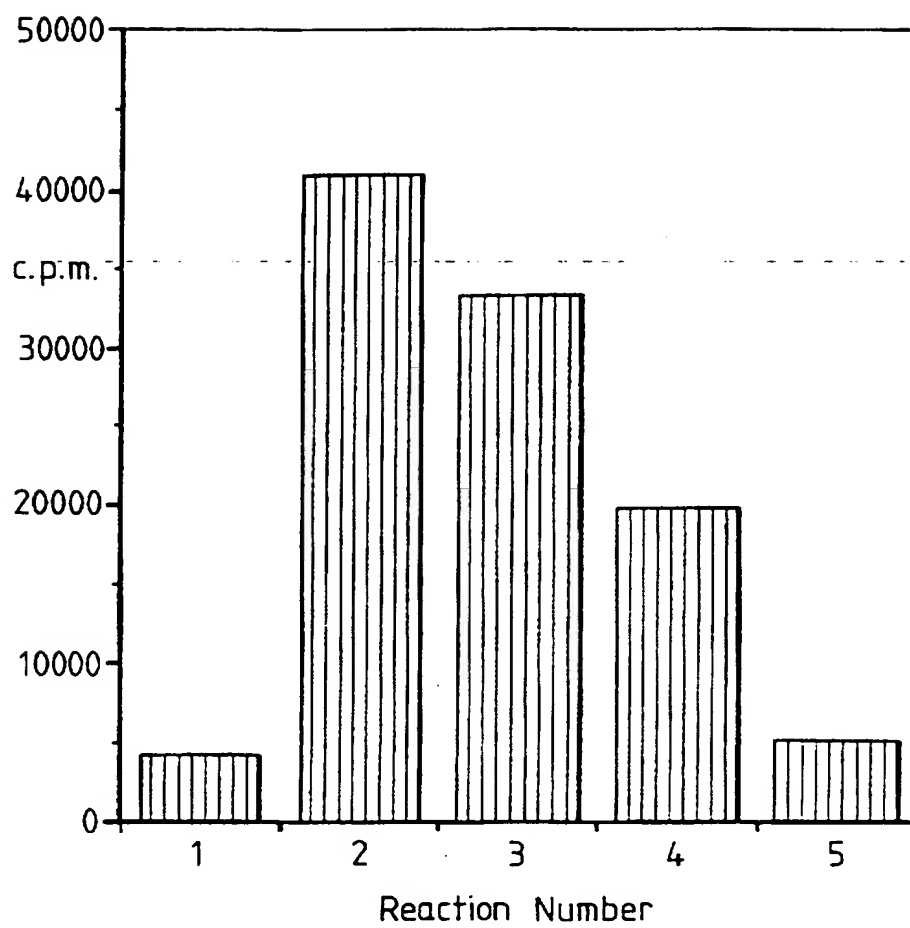
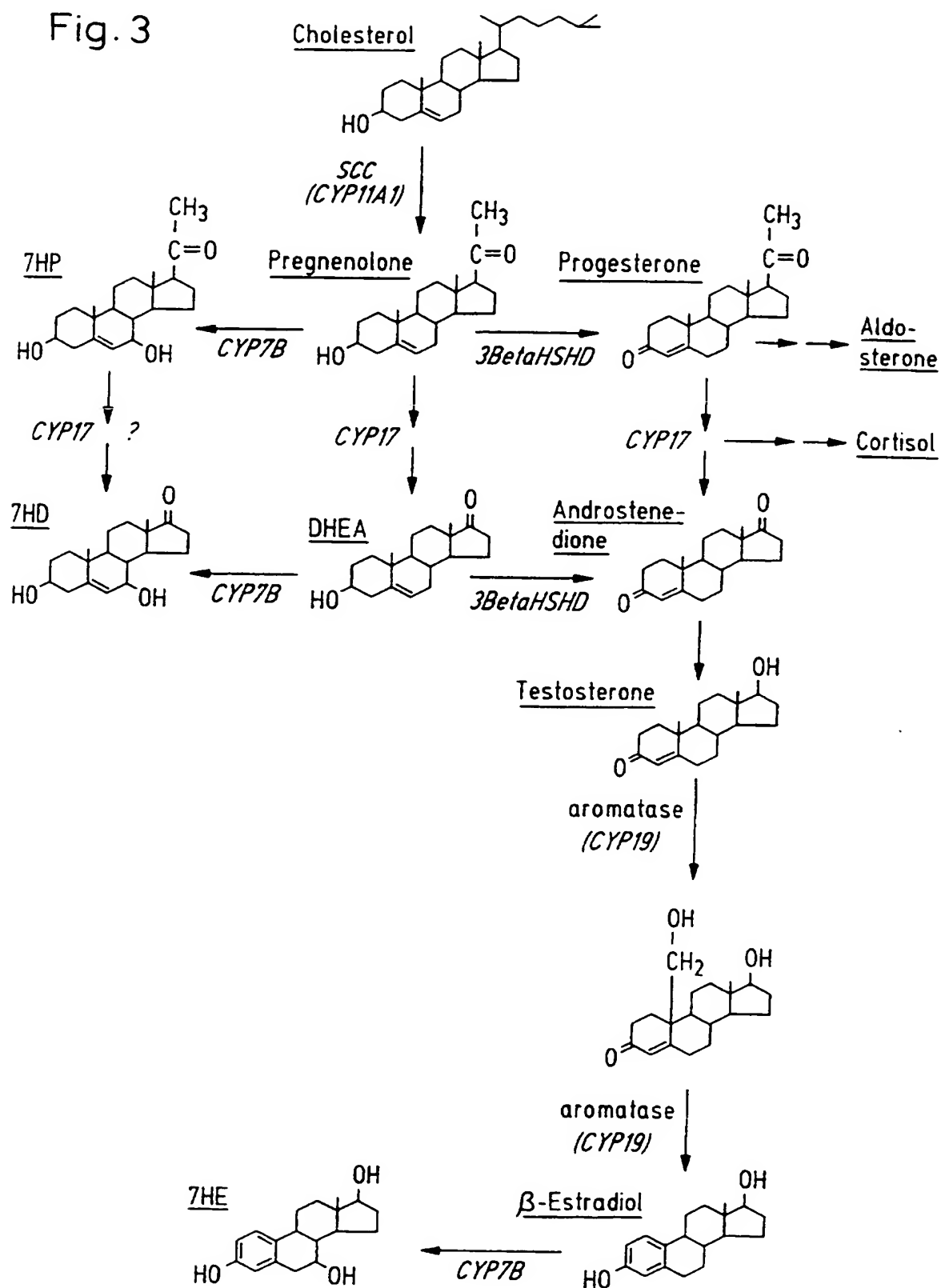


Fig. 2



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Fig. 3



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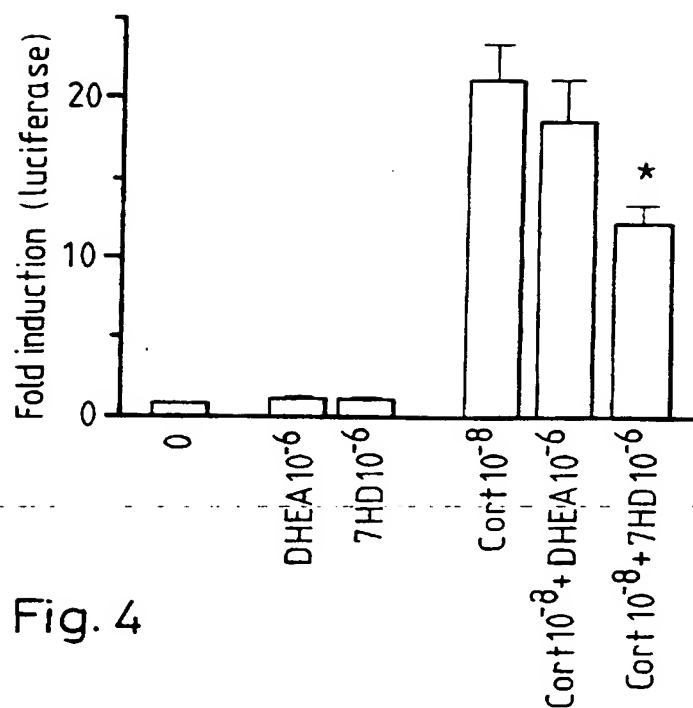


Fig. 4

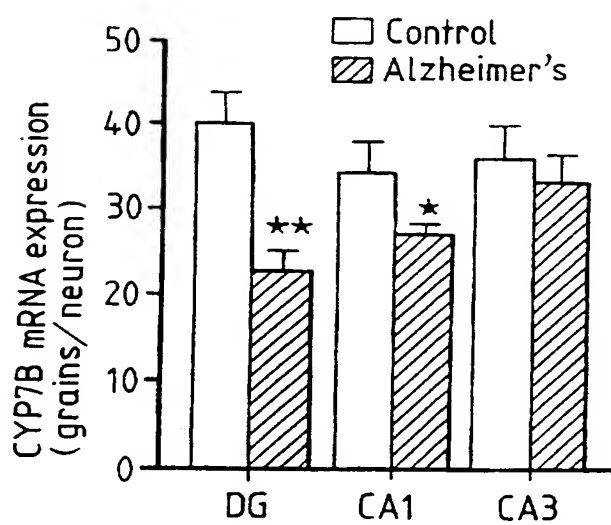
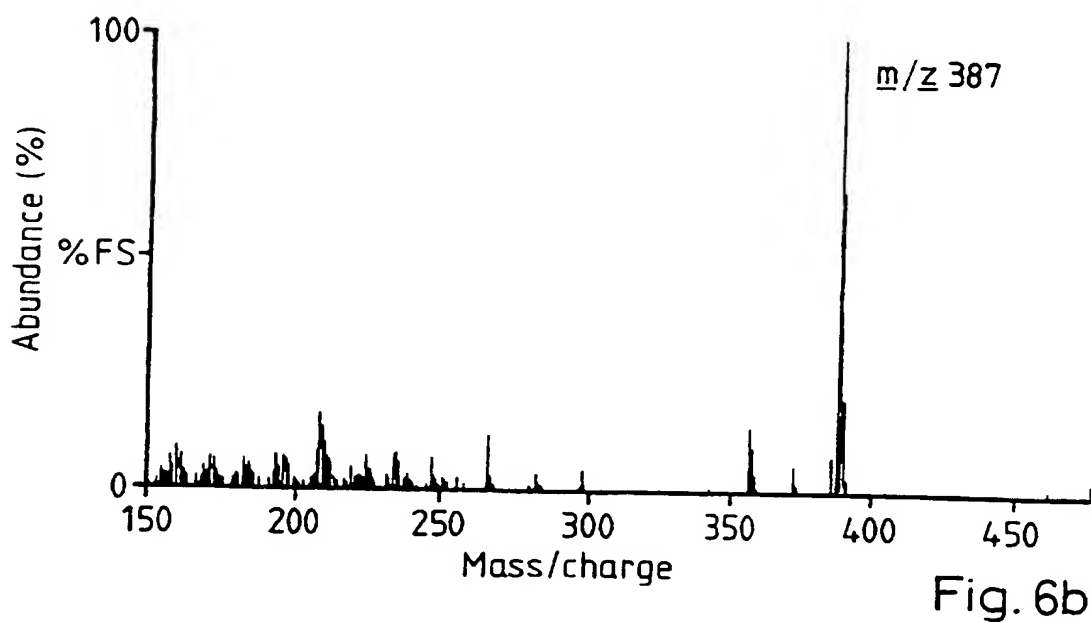
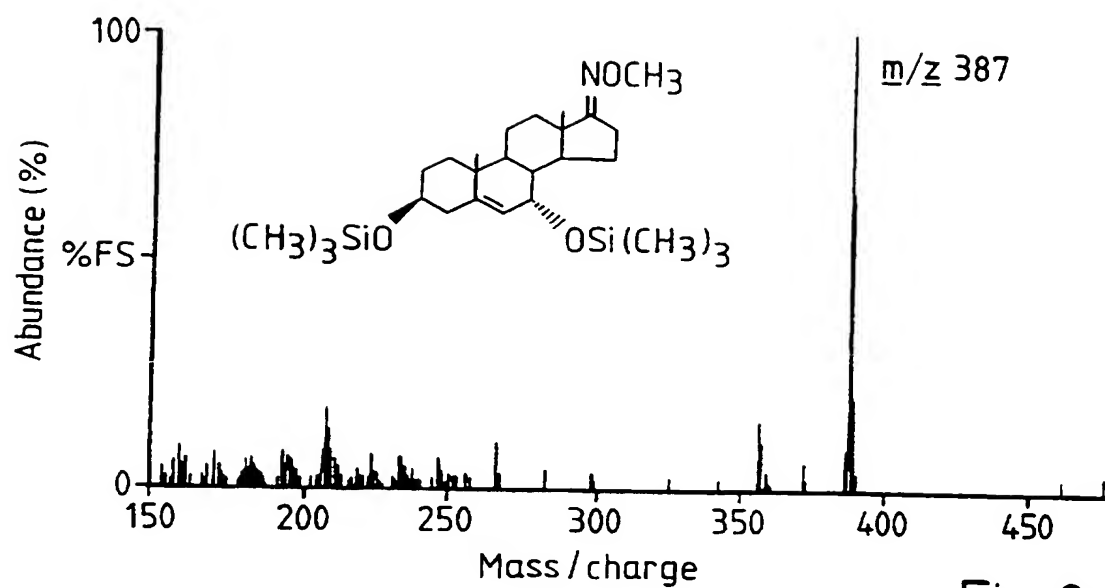


Fig. 5

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## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

|   |  |  |  |
|---|--|--|--|
| (51) International Patent Classification <sup>6</sup> :<br><b>A61K 31/565, C07J 1/00, C12P 33/00,<br/>A61K 39/395</b>   |  | <b>A3</b>  | (11) International Publication Number: <b>WO 97/37664</b><br>(43) International Publication Date: 16 October 1997 (16.10.97) |
| (21) International Application Number: <b>PCT/GB97/00955</b><br>(22) International Filing Date: 4 April 1997 (04.04.97)   |  | of Edinburgh, Western General Hospital, Edinburgh EH4 2XU (GB). LECKIE, Caroline, McKenzie [GB/GB]; Molecular Medicine Centre, Molecular Endocrinology, The University of Edinburgh, Western General Hospital, Edinburgh EH4 2XU (GB).   |  |
| (30) Priority Data:<br>9607289.7 9 April 1996 (09.04.96) GB<br>9608445.4 24 April 1996 (24.04.96) GB<br>9704905.0 10 March 1997 (10.03.97) GB   |  | (74) Agent: DOLAN, Anthony, Patrick; British Technology Group Ltd., Patents Dept., 101 Newington Causeway, London SE1 6BU (GB).  |  |
| (71) Applicant (for all designated States except US): <b>BRITISH TECHNOLOGY GROUP LTD. [GB/GB]; 101 Newington Causeway, London SE1 8BU (GB).</b>  |  | (81) Designated States: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, TJ, TM, TR, TT, UA, UG, US, UZ, VN, ARIPO patent (GH, KE, LS, MW, SD, SZ, UG), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG). |  |
| (72) Inventors; and<br>(75) Inventors/Applicants (for US only): <b>LATHE, Richard [GB/GB]; University of Edinburgh, West Mains Road, Edinburgh EH9 3JQ (GB). ROSE, Kenneth, Andrew [GB/GB]; University of Edinburgh, West Mains Road, Edinburgh EH9 3JQ (GB). SECKL, Jonathan, Robert [GB/GB]; Molecular Medicine Centre, Molecular Endocrinology, The University of Edinburgh, Western General Hospital, Edinburgh EH4 2XU (GB). BEST, Ruth [GB/GB]; Molecular Medicine Centre, Molecular Endocrinology, The University of Edinburgh, Western General Hospital, Edinburgh EH4 2XU (GB). YAU, Joyce, Lai, Wah [GB/GB]; Molecular Medicine Centre, Molecular Endocrinology, The University</b> |  | <b>Published</b><br><i>With international search report.</i><br><i>Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i>  |  |
|   |  | (88) Date of publication of the international search report: 4 December 1997 (04.12.97)  |  |
| (54) Title: <b>USE OF 7 ALPHA-SUBSTITUTED STEROIDS TO TREAT NEUROPSYCHIATRIC, IMMUNE OR ENDOCRINE DISORDERS</b>   |  |  |  |
| (57) Abstract<br><br>Use is provided for a 7 $\alpha$ -hydroxy or 7-oxo substituted 3 $\beta$ -hydroxy-steroid possessing the carbon skeleton of cholesterol, androster ne, pregnenolone or estradiol, or an analogue thereof substituted independently at one or both of the 7- and 3-positions with an ester or ether group, in the manufacture of a pharmaceutical composition for the therapy of neuropsychiatric, immune and/or endocrine disorders or for inducing cognitive enhancement. Uses for Cyp7b enzymes in producing such steroids is also provided together with various novel steroids and test kits and methods for diagnosing the disorders.                               |  |  |  |

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| BF | Burkina Faso             | GR | Greece                                   |    |  | TR | Turkey                   |
| BG | Bulgaria                 | HU | Hungary                                  | ML | Mali   | TT | Trinidad and Tobago      |
| BJ | Benin                    | IE | Ireland                                  | MN | Mongolia                                     | UA | Ukraine                  |
| BR | Brazil                   | IL | Israel                                   | MR | Mauritania                                   | UG | Uganda                   |
| BY | Belarus                  | IS | Iceland                                  | MW | Malawi                                       | US | United States of America |
| CA | Canada                   | IT | Italy                                    | MX | Mexico                                       | UZ | Uzbekistan               |
| CF | Central African Republic | JP | Japan                                    | NE | Niger  | VN | Viet Nam                 |
| CG | Congo                    | KE | Kenya                                    | NL | Netherlands                                  | YU | Yugoslavia               |
| CH | Switzerland              | KG | Kyrgyzstan                               | NO | Norway                                       | ZW | Zimbabwe                 |
| CI | Côte d'Ivoire            | KP | Democratic People's<br>Republic of Korea | NZ | New Zealand                                  |    |                          |
| CM | Cameroon                 |    |  | PL | Poland                                       |    |                          |
| CN | China                    | KR | Republic of Korea                        | PT | Portugal                                     |    |                          |
| CU | Cuba                     | KZ | Kazakhstan                               | RO | Romania                                      |    |                          |
| CZ | Czech Republic           | LC | Saint Lucia                              | RU | Russian Federation                           |    |                          |
| DE | Germany                  | LI | Liechtenstein                            | SD | Sudan  |    |                          |
| DK | Denmark                  | LK | Sri Lanka                                | SE | Sweden                                       |    |                          |
| EE | Estonia                  | LR | Liberia                                  | SG | Singapore                                    |    |                          |

## INTERNATIONAL SEARCH REPORT

International Application No

PC/GB 97/00955

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 6 A61K31/565 C07J1/00 C12P33/00 A61K39/395

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 A61K C07J C12P

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

| Category * | Citation of document, with indication, where appropriate, of the relevant passages   | Relevant to claim No. |
|------------|--|-----------------------|
| X          | WO 94 03176 A (HUMANETICS CORP.) 17<br>February 1994<br>see claims; examples<br>---  | 1-4,<br>28-33         |
| P,X        | WO 96 12810 A (UNIVERSITY OF EDINBURGH) 2<br>May 1996<br>cited in the application<br>see claim 20<br>---   | 7                     |
| P,X        | K.A. ROSE ET AL.: "Cyp7b, a novel brain<br>cytochrome P450, catalyzes the synthesis<br>of neurosteroids 7alpha-hydroxy<br>dehydroepiandrosterone and 7 alpha-hydroxy<br>pregnenolone."<br>BIOCHEMISTRY,<br>vol. 94, no. 10, 1997,<br>pages 4925-4930, XP002042014<br>see the whole document<br>--- | 11,12,<br>18-21       |

-/--



Further documents are listed in the continuation of box C.



Patent family members are listed in annex.

## \* Special categories of cited documents:

\*A\* document defining the general state of the art which is not considered to be of particular relevance

\*E\* earlier document but published on or after the international filing date

\*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

\*O\* document referring to an oral disclosure, use, exhibition or other means

\*P\* document published prior to the international filing date but later than the priority date claimed

\*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

\*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

\*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

\*Z\* document member of the same patent family

Date of the actual completion of the international search

29 September 1997

Date of mailing of the international search report

17.10.97

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2

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Fax: (+31-70) 340-3016

Authorized officer

Klaver, T

# INTERNATIONAL SEARCH REPORT

nal Application No

PCT/GB 97/00955

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

| Category * | Citation of document, with indication, where appropriate, of the relevant passages  | Relevant to claim No. |
|------------|---|-----------------------|
| P,X        | K.A. ROSE ET AL.: "Steroid modification in brain: hydroxylation of pregnenolone and DHEA by the novel cytochrome P450, Cyp7b."<br>J. ENDOCRINOL.,<br>vol. 152, 1997,<br>page P280 XP002042015<br>see the whole document | 11,12,<br>18-21       |
| A          | ---<br>G. STAPLETON ET AL.: "A novel cytochrome P450 expressed primarily in brain."<br>J. BIOL. CHEM.,<br>vol. 270, no. 50, 1995,<br>pages 29739-29745, XP002042016<br>see the whole document                           |                       |
| A          | ---<br>EP 0 648 842 A (NORTHEASTERN OHIO UNIVERSITIES) 19 April 1995<br>-----   |                       |



# INTERNATIONAL SEARCH REPORT

International application No.

GB 97/ 00955

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:
  
2. ☒ Claims Nos.: 1, 28  
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:  
It is not clear which compounds are described by "...or a derivative thereof..." nor is it clear which diseases are meant by descriptions like "...neuro psychiatric, immune and/or endocrine disorders...". The search has therefore been limited to the examples mentioned in the claims and/or description.
  
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

See annex

1. ☒ As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
  
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
  
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
  
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☒ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

# INTERNATIONAL SEARCH REPORT

International Application No. PCT/GB 97 00955

FURTHER INFORMATION CONTINUED FROM PCT/ISA210

- 1) Claims 1-4, 28-33: Use of 7-substituted 3 $\beta$ -hydroxy steroids and novel steroids, for the treatment of neuropsychiatric, immune and/or endocrine disorders.
- 2) Claims 5 - 2 : Use of Cyp7b to manufacture assay kits, to produce 7-hydroxy steroids, antibodies, and targeted drugs for gene therapy.
- 3) Claims 23 - 27 : Novel steroids of formula 1a and 1b.

# INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/GB 97/00955

| Patent document<br>cited in search report | Publication<br>date | Patent family<br>member(s) | Publication<br>date |
|---|---------------------|----------------------------|---------------------|
| WO 9403176 A                              | 17-02-94            | US 5292730 A               | 08-03-94            |
|   |                     | AU 4997093 A               | 03-03-94            |
|   |                     | CA 2141436 A               | 17-02-94            |
|   |                     | EP 0746322 A               | 11-12-96            |
|   |                     | JP 8505602 T               | 18-06-96            |
|   |                     | US 5585371 A               | 17-12-96            |
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| WO 9612810 A                              | 02-05-96            | AU 3670395 A               | 15-05-96            |
|   |                     | EP 0795017 A               | 17-09-97            |
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| EP 648842 A                               | 19-04-95            | US 5420028 A               | 30-05-95            |
|   |                     | EP 0648840 A               | 19-04-95            |
|   |                     | JP 7284393 A               | 31-10-95            |
|   |                     | JP 7284388 A               | 31-10-95            |
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